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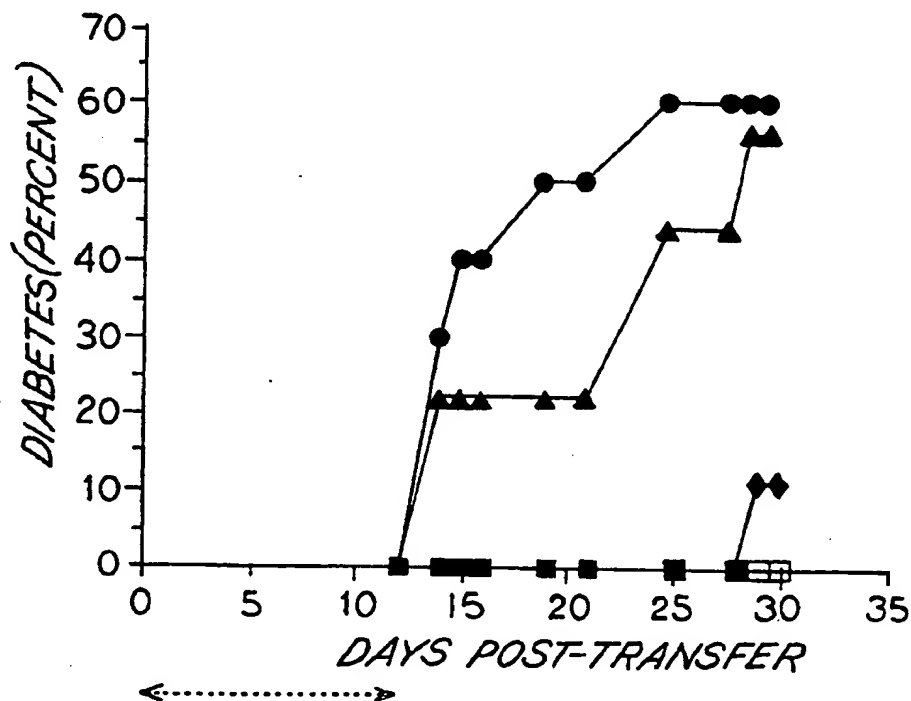
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(57) Abstract

A method for the prevention of insulin dependent (type I) diabetes. The method comprises administration of an antibody, polypeptide or other molecule recognizing VLA4.



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TREATMENT FOR INSULIN DEPENDENT DIABETES

This application is a continuation-in-part of U.S. Patent Application Serial No. 08/029,330, filed February 9, 1993.

FIELD OF THE INVENTION

The present invention relates to a treatment for insulin dependent (type-I) diabetes. More particularly, this invention relates to the use of antibodies recognizing the integrin VLA4 (very late antigen 4) in the prevention of diabetes.

BACKGROUND OF THE INVENTION

Insulin dependent diabetes (also termed type-I diabetes and formerly juvenile onset diabetes mellitus) has been classified during the past two decades as a chronic autoimmune disease. In this disorder, cells producing insulin (β cells) within the pancreatic islets are selectively targeted and destroyed by a cellular infiltrate of the pancreas. This inflammatory infiltrate affecting the islets has been termed insulinitis. Cells producing insulin comprise the majority of islet cells but less than 2% of the total pancreatic mass (Castano and Eisenbarth, 1990, [1]; Fujita et al., 1982 [2]; Foulis et al., 1986 [3]). The development of type I diabetes can conceptually be divided into six stages, beginning with genetic susceptibility and ending with complete β cell destruction (Eisenbarth, 1986 [4]). Stage I is genetic susceptibility, which is a necessary but insufficient condition for development of the disease. A hypothetical triggering event (Stage II) leads to active autoimmunity against β cells (Stage III). In Stage III, the β cell mass is hypothesized to decline

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and immunologic abnormalities such as autoantibodies directed against insulin and islet cytoplasmic antigens are found. Stimulated insulin secretion is still preserved at this stage. Over a period of years, however, the progressive loss of β cells leads to diminished insulin secretion with intravenous glucose tolerance tests (IVGTT) while the individual is still normoglycemic (Stage IV). Overt diabetes (i.e., diabetes onset or clinical manifestation of disease characterized by hyperglycemia) is Stage V, and can develop years later when approximately 90% of pancreatic β cells are destroyed. In Stage V when overt diabetes is first recognized, some residual insulin production remains (as demonstrated by the presence of the connecting peptide of proinsulin, C peptide, in the serum) but the individual usually requires exogenous insulin for life. Finally, in Stage VI, even the remaining β cells are destroyed and C peptide can no longer be detected in the circulation.

While the initiating factor(s) and specific sequence of events leading to diabetes, including the relative importance of different cell types and cytokines, are still widely debated, a key role is generally recognized for self-antigen reactive T cells (Miller et al., 1988 [5]; Harada and Makino, 1986 [6]; Koike et al., 1987 [7]; Makino et al., 1986 [8]). In addition to T lymphocytes, insulinitis is characterized by macrophages, dendritic cells (Voorbij et al., 1989 [9]) and B cells, which may serve as professional antigen presenting cells (APC). Macrophages may also destroy islet β cells themselves by release of cytokines or free radicals (Nomikos et al., 1986 [10]). Thus, autoimmune diabetes relies upon both cellular migration and immune stimulation of newly resident cells.

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Cell trafficking to inflammatory sites is regulated by accessory molecules LFA-1, MAC-1 and VLA4 (Larson and Springer, 1990 [11]; Hemler et al., 1990 [12]) on the surface of lymphocytes (LFA-1, VLA4) and macrophages (Mac-1, VLA4), and by their counter-ligands ICAM (for LFA-1 and MAC-1), and VCAM (for VLA4) which are unregulated by cytokines on vascular endothelium (Larson and Springer, 1990 [11]; Lobb, 1992 [13]; Osborn, 1990, [14]). In addition, VLA4 binds to an extracellular matrix component, the CS-1 domain of fibronectin (FN) (Wayner et al., 1989 [15]). The relative importance of these pathways, for example, LFA-1 and VLA4 on lymphocytes or MAC-1 and VLA4 on monocytes, in controlling cell migration is still a subject of investigation. In vitro data suggest that the differential use of these pathways appears to depend upon the activation status of both the leukocytes and endothelial cells (Shimizu et al., 1991 [16]). Their ability to control cell migration to inflammatory sites in vivo has been directly demonstrated with monoclonal antibodies (mAbs) to ICAM, MAC-1 or VLA4 inhibiting various animal models of disease (Barton et al., 1989 [17], phorbol ester-induced rabbit lung inflammation; Issekutz and Issekutz, 1991 [18], delayed type hypersensitivity; Issekutz, 1991 [19], adjuvant-induced arthritis; Yednock et al., 1992 [20], transfer of experimental allergic encephalomyelitis (EAE); Lobb, 1992 [21], asthma).

ICAM and VCAM are also found on the surface of macrophages and dendritic cells in lymphoid tissues (Dustin et al., 1986 [22]; Rice et al., 1990 [23]; Rice et al., 1991 [24]). Their distribution on these professional APC is consistent with functional data indicating a role for LFA-1 and VLA4 in T cell

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activation (Shimizu et al., 1990 [25], Burkly et al., 1991 [26]). However, numerous other receptor-ligand pairs including CD4/ MHC class II and CD8/MHC class I (Rudd et al., 1989 [27]), CD2/LFA-3 (Moingeon et al., 1989 [28]), CD28/B7 (Harding et al., 1992 [29]) may also support adhesion or costimulate T cells during T/APC or T/target cell interactions. The specific contributions of these numerous pathways in the development of diabetes is unresolved. Because there are multiple molecular pathways for cell adhesion and T cell activation, it is not possible to predict whether intervention in one or more of these pathways might affect onset or severity of diabetes disease, and, in particular, which of these pathways are crucial or relevant to the disease process.

Antibodies directed to T cells have been utilized in murine and rat models for spontaneous diabetes and adoptive transfer of diabetes to deplete T cells and thus prevent disease (see, e.g., Harada and Makino, 1986 [6], anti-Thy 1.2; Koike et al., 1987 [7], Miller et al., 1988 [5] and Shizuru et al., 1988 [30], anti-CD4; Barlow and Like, 1992 [31], anti-CD2; Like et al., 1986 [32], anti-CD5 and anti-CD8). In addition, an antibody directed to the complement receptor type 3 (CR3) molecule or MAC-1 on macrophages has been utilized to prevent macrophage and T cell infiltration of pancreatic tissue in a murine adoptive transfer model of disease (Hutchings et al., 1990 [33]). It is unknown whether VLA4 is relevant to insulinitis or to the activity of islet-specific cells after localization in the pancreas.

Current treatment protocols suggested for type I diabetes have included certain immunomodulatory drugs summarized by Federlin and Becker [34] and references cited therein. A long prediabetic period with

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immunologic abnormalities and progressive β cell destruction suggests it may be possible to halt β cell loss with immune intervention (Castano and Eisenbarth, 1990 [1]).

5 Suggested agents/protocols have included certain immunomodulatory and immunosuppressive agents: levamisole, theophyllin, thymic hormones, ciamexone, anti-thymocyte globulin, interferon, nicotinamide, gamma globulin infusion, plasmapheresis or white cell
10 transfusion. Agents such as cyclosporin A and azathioprine which impair T cell activation and T cell development, respectively, have been used in clinical trials (Zielasek et al., 1989 [35]). The most
15 promising results have been achieved with cyclosporin A (Castano and Eisenbarth, 1990 [1]). Federlin and Becker, 1990 [34] suggest, however, that cyclosporin A may not be recommended for general or long-term use because of toxic side effects, at least when given in
20 higher doses. Higher doses of cyclosporin, or in combination with other immunosuppressive drugs, or both, have been associated with the development of lymphoma and irreversible kidney damage (Eisenbarth, 1986 [4]; Eisenbarth, 1987 [36]). Additional studies
25 on other suggested agents are necessary to assess safety and efficacy. Even the cyclosporin A studies show that its efficacy in maintaining remission of diabetes is for one year in about 30-60% of new onset diabetes. Within 3 years, however, remissions are almost invariably lost (Castano and Eisenbarth, 1990
30 [1]). Treatment protocols after onset of disease are particularly problematic, since, for example, at the time diabetes is diagnosed in humans, insulinitis has typically progressed already to a loss of more than 80% of the β cells. Thus, it is possible that cyclosporin
35 A may be preventing further β cell destruction, but so

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few β cells may be present at the onset of the diabetes that they cannot maintain a non-diabetic state over time (Castano and Eisenbarth, 1990 [1]). Suppression of insulinitis and/or prevention of disease may be more successful if the treatment could start at an earlier phase, i.e., before disease onset.

There are two major prerequisites in order to develop any preventative treatment for diabetes disease: (1) the ability to accurately identify the prediabetic individual and (2) the development of safe, specific and effective preventive treatments. Significant progress has been made in identifying prediabetic individuals, however, much work remains in the development of safe, specific and effective preventive treatments as discussed and reviewed by Eisenbarth and colleagues (see, e.g., Ziegler and Eisenbarth, 1990 [37]; Ziegler et al., 1990 [38]; Ziegler et al., 1990 [39]). It has been possible to identify certain risk factors and at-risk groups for type I diabetes and thus to predict individuals most likely to go on to clinical disease and to estimate the approximate rate of disease onset in these individuals. The ability to identify individuals with susceptibility to diabetes or to predict type I diabetes in the pre-clinical stage by the combination of genetic (HLA typing), immunological (islet and insulin autoantibodies) and metabolic (first phase insulin secretion to intravenous glucose preceding the development of hyperglycemia) markers makes the identification and use of prophylactic immunotherapeutic drugs and protocols possible during the evolution of the autoimmune disease process when β cell destruction is only partial. To date, there has been little success, however, in treating human diabetes. Generally, because human treatment has been

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us d only after onset of the disease, treatment was
f llowed by a temporary complet or partial remission
only in a certain number f patients. Sinc
5 immunosuppressive mechanisms may prevent insulitis
and/or diabetes, there is a need for immunosuppressive
components for use in the prediabetic stage. In
particular, there is a need for safer and more
specifically acting compounds, e.g., monoclonal
antibodies, which inhibit entry of effector cells into
10 the pancreas or function of those cell which may have
already entered the islets of Langerhans.

It has now been surprisingly discovered that
administering an anti-VLA4 antibody significantly
reduced the incidence of diabetes, in a rodent model of
15 diabetes disease. The NOD mouse model of diabetes is a
well established model directly comparable to human
type-I diabetes. Using an adoptively transferred
disease experimental protocol, irradiated non-diabetic
NOD mice were administered splenocytes from
20 spontaneously diabetic NOD mice for the acute transfer
of the disease. These splenocytes were treated with
anti-VLA4 antibody before administration and the
recipients were also treated for various periods of
time after the transfer with anti-VLA4 antibody.

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SUMMARY OF THE INVENTION

Accordingly, the present invention provides novel methods for the treatment of insulin dependent (type I) diabetes in a prediabetic. In particular, the present invention provides a method for the prevention of insulin dependent diabetes comprising the step of administering to a prediabetic individual an anti-VLA4 antibody, such as antibody HP1/2 or a humanized anti-VLA4 antibody derived from HP1/2. Also contemplated is the use of analogous antibodies, antibody fragments, soluble proteins and small molecules that mimic the action of anti-VLA4 antibodies in the treatment of diabetes. In addition, the present invention provides a method for the treatment of diabetes by administering to a mammal, including a human, with a susceptibility to diabetes an antibody capable of binding to the $\alpha 4$ subunit of VLA4 in an amount effective to provide inhibition of the onset of diabetes. Also contemplated is the use of recombinant and chimeric antibodies, fragments of such antibodies, polypeptides or small molecules capable of binding $\alpha 4$ /VLA4. Also contemplated are soluble forms of the natural binding proteins for VLA 4, including soluble VCAM-1, VCAM-1 peptides or VCAM-1 fusion proteins as well as fibronectin, fibronectin having an alternatively spliced non-type III connecting segment and fibronectin peptides containing the amino acid sequence EILDV or a similar conservatively substituted amino acid sequence. These agents will act by competing with the cell-surface binding protein for VLA4.

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BRIEF DESCRIPTION OF THE DRAWINGS

Figur 1 is a graph depicting th effect of anti-VLA4 antib dy (R1-2) and contr ls on prevention of diabetes after adoptive transfer of spleen cells; the frequency of recipients which became diabetic and day of disease onset are shown for transfer of 2×10^7 splenocytes from diabetic (D) NOD donors without treatment (closed circles), with a non-specific rat IgG2b treatment (closed triangles), and with R1-2 anti-VLA4 treatment (closed diamonds), as well as for transfer of splenocytes from nondiabetic (Y) NOD donors (open squares); the splenocytes were transferred with R1-2 or rat IgG2b or without mAb, and then R1-2 or rat IgG2b was injected every other day through day 12 post transfer (n=8-10 for all groups).

Figure 2 is a graph depicting the effect of anti-VLA4 antibody (R1-2) and controls on prevention of diabetes after adoptive transfer of spleen cells; the frequency of recipients which became diabetic and day of disease onset are shown for transfer of 3×10^7 splenocytes from diabetic (D) NOD donors without treatment (closed circles), with a non-specific rat IgG2b treatment (closed triangles), and with R1-2 anti-VLA4 treatment (closed diamonds), as well as for transfer of splenocytes from nondiabetic (Y) NOD donors (open squares); the splenocytes were transferred with R1-2 or rat IgG2b or without mAb, and then R1-2 or rat IgG2b was injected every 3.5 days through day 25 post transfer (n=4-5 for all groups).

Figure 3 is a graph depicting the effect of anti-VLA4 antibody (R1-2) and controls on prevention of diabetes after adoptive transfer of spleen cells; the frequency of recipients which became diabetic and day of disease onset are shown for transfer of $2-3 \times 10^7$ splenocytes from diabetic (D) NOD donors without

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treatment (closed circles), with a non-specific rat IgG2b treatment (closed triangles), and with R1-2 anti-VLA4 treatment (closed diamonds), as well as for transfer of splenocytes from nondiabetic (Y) NOD donors (open squares) or for PBS alone (open circles); the splenocytes were transferred with R1-2 or rat IgG2b or without mAb, and then R1-2 or rat IgG2b was injected every 3.5 days through day 25 post transfer (n=5 for all groups).

Figure 4 is a bar graph depicting the effect of anti-VLA4 antibody (R1-2) and controls on the degree of insulinitis after adoptive transfer of spleen cells; the frequency of uninfiltrated islets (Grade 0-I infiltrate, stippled bar) and infiltrated islets (Grade II-IV insulinitis, solid bar) were quantitated and shown after transfer of cells treated with R1-2, rat IgG2b or without mAb, and then R1-2 or rat IgG2b injected every 3.5 days through day 25 with mice sacrificed when diabetic or on day 26 post-transfer. Pancreatic sections from n=4-5 mice were scored for each experimental group, i.e., Y→Y (non-diabetic donor cells) or D→Y (diabetic donor cells) into non-diabetic (Y) recipients with no mAb treatment, treatment with rat IgG2b or treatment with R1-2.

Figure 5 is a bar graph depicting the effect of anti-VLA4 antibody (R1-2) and controls on the degree of insulinitis after adoptive transfer of spleen cells; the frequency of uninfiltrated islets (Grade 0-I infiltrate, stippled bar) and infiltrated islets (Grade II-IV insulinitis, solid bar) were quantitated and shown after transfer of cells treated with R1-2, rat IgG2b or without mAb, and then R1-2 or rat IgG2b injected every other day through day 12 post-transfer, then maintained without further mAb injection until sacrificed when diabetic or on day 29 post-transfer. Pancreatic

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s cells from n=4-5 mice were scored for each experimental group, i.e., Y→Y (non-diabetic donor cells) or D→Y (diabetic donor cells) into non-diabetic (Y) recipients with no mAb treatment, treatment with rat IgG2b or treatment with R1-2.

Figure 6 is a graph depicting the effect of anti-VLA4 antibody (R1-2) and controls on prevention of diabetes in a spontaneous disease model for diabetes; the frequency of recipients which became diabetic and day of disease onset are shown for NOD mice without treatment (closed squares), with a non-specific rat IgG2b treatment (closed circles), and with R1-2 anti-VLA4 treatment (closed triangles); R1-2 or rat IgG2b was injected for 8 weeks in NOD mice twice weekly from week four to week twelve of age.

Figure 7 is a graph depicting the effect of VCAM 2D-IgG fusion protein and controls on prevention of diabetes after adoptive transfer of spleen cells; the frequency of recipients which became diabetic and day of disease onset are shown for transfer of 2×10^7 splenocytes from diabetic (D) NOD donors with an irrelevant rat LFA-3Ig fusion protein treatment (closed squares), and with VCAM 2D-IgG treatment (open circles) or of recipients which received PBS alone without cells transferred (closed triangles); the splenocytes were transferred with VCAM 2D-IgG or rat LFA-3Ig, and then VCAM 2D-IgG or rat LFA-3Ig was injected every other day through day 17 post-transfer (n = 5 for all groups).

Figure 8 is a schematic depicting structure of VCAM 2DIgG fusion protein described in Example 5. VCAM 2D-IgG is a soluble form of the ligand for VLA4 (VCAM1) and consists of the two N-terminal domains of VCAM1 fused to the human IgG1 heavy chain constant region sequences (Hinges, C_H2 and C_H3).

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DETAILED DESCRIPTION OF THE INVENTION

Th invention relates to a treatment including the prevention of insulin dependent (type I) diabetes. More particularly, the invention relates to the use of antibodies to VLA4 in the treatment of diabetes in a prediabetic individual. The term "prediabetic" is intended to mean an individual at risk for the development of diabetes disease (e.g., genetically predisposed) at any stage in the disease process prior to overt diabetes or diabetes onset. The term "diabetic" is intended to mean an individual with overt hyperglycemia (i.e., fasting blood glucose levels \geq 250 mg/dL). The term "overt diabetes" or "diabetes onset" is intended to mean a disease state in which the pancreatic islet cells are destroyed and which is manifested clinically by overt hyperglycemia (i.e., fasting blood glucose levels \geq 250 mg/dL).

In the first aspect, the invention provides a method of treatment of diabetes comprising the step of administering a composition capable of binding to, including blocking or coating, the VLA4 antigens on the surface of VLA4-positive cells, including lymphocytes and macrophages. For purposes of the invention, the term "binding to VLA4 antigens" is intended to mean reacting with VLA4 antigens on cells and thereby interfering with interactions between VLA4 antigens and either VCAM-1 or fibronectin on the surface of other cells or thereby inducing a change in the function of the VLA4-positive cells. As demonstrated herein, such binding, including blocking or coating, of VLA4 antigens results in a prevention in or protection against the incidence of diabetes. This demonstration utilized a monoclonal antibody against VLA4 as a binding agent which effectively blocked or coated the VLA4 antigens. Those skilled in the art will recognize

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that, given this demonstration, any agent that can bind to, including those that can block or coat, VLA4 antigens can be successfully used in the method of the invention. Thus, for purposes of the invention, any agent capable of binding to VLA4 antigens on the surface of VLA4-bearing cells and which may effectively block or coat VLA4 antigens, is considered to be an equivalent of the monoclonal antibody used in the examples herein. For example, the invention contemplates as binding equivalents at least peptides, peptide mimetics, carbohydrates and small molecules capable of binding VLA4 antigens on the surface of VLA4-bearing cells.

In a preferred embodiment, the agent that is used in the method of the invention to bind to, including block or coat, cell-surface VLA4 antigens is a monoclonal antibody or antibody derivative. Preferred antibody derivatives for treatment, in particular for human treatment, include humanized recombinant antibodies, chimeric recombinant antibodies, Fab, Fab', F(ab')₂ and F(v) antibody fragments, and monomers or dimers of antibody heavy or light chains or intermixtures thereof. Thus, monoclonal antibodies against VLA4 are a preferred binding agent in the method according to the invention.

The technology for producing monoclonal antibodies is well known. Briefly, an immortal cell line (typically myeloma cells) is fused to lymphocytes (typically splenocytes) from a mammal immunized with whole cells expressing a given antigen, e.g., VLA4, and the culture supernatants of the resulting hybridoma cells are screened for antibodies against the antigen. (See, generally, Kohler et al., 1975 [40]).

Immunization may be accomplished using standard procedures. The unit dose and immunization regimen

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depend on the species of mammal immunized, its immune status, the body weight of the mammal, etc. Typically, the immunized mammals are bled and the serum from each blood sample is assayed for particular antibodies using appropriate screening assays. For example, anti-VLA4 antibodies may be identified by immunoprecipitation of ^{125}I -labeled cell lysates from VLA4-expressing cells. (See, Sanchez-Madrid et al. 1986 [41] and Hemler et al. 1987 [42]). Anti-VLA4 antibodies may also be identified by flow cytometry, e.g., by measuring fluorescent staining of Ramos cells incubated with an antibody believed to recognize VLA4 (see, Elices et al., (1990) [43]). The lymphocytes used in the production of hybridoma cells typically are isolated from immunized mammals whose sera have already tested positive for the presence of anti-VLA4 antibodies using such screening assays.

Typically, the immortal cell line (e.g., a myeloma cell line) is derived from the same mammalian species as the lymphocytes. Preferred immortal cell lines are mouse myeloma cell lines that are sensitive to culture medium containing hypoxanthine, aminopterin and thymidine ("HAT medium").

Typically, HAT-sensitive mouse myeloma cells are fused to mouse splenocytes using 1500 molecular weight polyethylene glycol ("PEG 1500"). Hybridoma cells resulting from the fusion are then selected using HAT medium, which kills unfused and unproductively fused myeloma cells (unfused splenocytes die after several days because they are not transformed). Hybridomas producing a desired antibody are detected by screening the hybridoma culture supernatants. For example, hybridomas prepared to produce anti-VLA4 antibodies may be screened by testing the hybridoma culture supernatant for secreted antibodies having the ability

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t bind to a recombinant α_4 -subunit-expressing cell line, such as transfected K-562 cells (see, Ellis et al. [43]).

5 To produce anti-VLA4 antibodies, hybridoma cells that tested positive in such screening assays were cultured in a nutrient medium under conditions and for a time sufficient to allow the hybridoma cells to secrete the monoclonal antibodies into the culture medium. Tissue culture techniques and culture media
10 suitable for hybridoma cells are well known. The conditioned hybridoma culture supernatant may be collected and the anti-VLA4 antibodies optionally further purified by well-known methods.

Alternatively, the desired antibody may be
15 produced by injecting the hybridoma cells into the peritoneal cavity of an unimmunized mouse. The hybridoma cells proliferate in the peritoneal cavity, secreting the antibody which accumulates as ascites fluid. The antibody may be harvested by withdrawing
20 the ascites fluid from the peritoneal cavity with a syringe.

Several mouse anti-VLA4 monoclonal antibodies have been previously described (see, e.g., Sanchez-Madrid et al., 1986 [41]; Hemler et al., 1987 [42]; Pulido et
25 al., 1991 [44]). These anti-VLA4 monoclonal antibodies such as HP1/2 and other anti-VLA4 antibodies (e.g., mAb HP2/1, HP2/4, L25, P4C2, P4G9) capable of recognizing the α chain of VLA4 will be useful in the methods of treatment according to the present invention. Anti-
30 VLA4 antibodies that will recognize the VLA- α_4 chain epitopes involved in binding to VCAM-1 and fibronectin ligands (i.e., antibodies which can bind to VLA4 at a site involved in ligand recognition and block VCAM-1 and fibronectin binding) are preferred. Such
35 antibodies have been defined as B epitope-specific

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antibodies (B1 or B2) (see, Pulido et al. (1991) [36]) and are preferred anti-VLA4 antibodies according to the present invention. The R1-2 antibody used as described herein is a B epitope type antibody.

5 Human monoclonal antibodies against VLA4 are another preferred binding agent which may block or coat VLA4 antigens in the method of the invention. These may be prepared using *in vitro*-primed human splenocytes, as described by Boerner et al., 1991 [45].
10 Alternatively, they may be prepared by repertoire cloning as described by Persson et al., 1991 [46] or by Huang and Stollar, 1991 [47]. Another preferred binding agent which may block or coat VLA4 antigens in the method of the invention is a chimeric recombinant
15 antibody having anti-VLA4 specificity and a human antibody constant region. Yet another preferred binding agent which may block or coat VLA4 antigens in the method of the invention is a humanized recombinant antibody having anti-VLA4 specificity. Humanized
20 antibodies may be prepared, as exemplified in Jones et al., 1986 [48]; Riechmann, 1988, [49]; Queen et al., 1989 [50]; and Orlandi et al., 1989 [51]. Preferred binding agents including chimeric recombinant and humanized recombinant antibodies with B epitope
25 specificity have been prepared and are described in co-pending and co-assigned U.S. Patent Application Serial No. 08/004,798, filed January 12, 1993 [52]. The starting material for the preparation of chimeric (mouse V - human C) and humanized anti-VLA4 antibodies
30 may be a murine monoclonal anti-VLA4 antibody as previously described, a monoclonal anti-VLA4 antibody commercially available (e.g., HP2/1, Amac International, Inc., Westbrook, Maine), or a monoclonal anti-VLA4 antibody prepared in accordance with the
35 teaching herein. For example, the variable regions of

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the heavy and light chains of the anti-VLA4 antibody HP1/2 have been cloned, sequenced and expressed in combination with constant regions of human immunoglobulin heavy and light chains. Such a chimeric HP1/2 antibody is similar in specificity and potency to the murine HP1/2 antibody, and may be useful in methods of treatment according to the present invention. The HP1/2 V_H DNA sequence and its translated amino acid sequences are set forth in SEQ ID NO: 1 and SEQ ID NO: 2, respectively. The HP1/2 V_K DNA sequence and its translated amino acid sequence are set forth in SEQ ID NO: 3 and SEQ ID NO: 4, respectively. Similarly, humanized recombinant anti-VLA4 antibodies may be useful in these methods. A preferred humanized recombinant anti-VLA4 antibody is an AS/SVMDY antibody, for example, the AS/SVMDY antibody produced by the cell line deposited with the ATCC on November 3, 1992 and given accession no. CRL 11175. The AS/SVMDY humanized antibody is at least equipotent with or perhaps more potent than the murine HP1/2 antibody. The AS V_H DNA sequence and its translated amino acid sequences are set forth in SEQ ID NO: 5 and SEQ ID NO: 6, respectively. The SVMDY V_K DNA sequence and its translated amino acid sequence are set forth in SEQ ID NO: 7 and SEQ ID NO: 8, respectively.

Those skilled in the art will recognize that any of the above-identified antibody or antibody derivative binding agents can also act in the method of the invention by binding to the receptor for VLA4, and may block or coat the cell-surface VLA4 antigen. Thus, antibody and antibody derivative binding agents according to the invention may include embodiments having binding specificity for VCAM-1 or fibronectin, since these molecules appear to either be important in the adhesion cells or the extracellular matrix or

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interfere with traffic of cells through tissues and blood.

Alternatively, the binding agents used in the method according to the invention may not be antibodies or antibody derivatives, but rather may be soluble forms of the natural binding proteins for VLA4. These binding agents include soluble VCAM-1, VCAM-1 peptides, or VCAM-1 fusion proteins as well as fibronectin, fibronectin having an alternatively spliced non-type III connecting segment and fibronectin peptides containing the amino acid sequence EILDV or a similar conservatively substituted amino acid sequence. These binding agents will act by competing with the cell-surface binding protein for VLA4.

In this method according to the first aspect of the invention, VLA4 binding agents are preferably administered parenterally. The VLA4 binding agents are preferably administered as a sterile pharmaceutical composition containing a pharmaceutically acceptable carrier, which may be any of the numerous well known carriers, such as water, saline, phosphate buffered saline, dextrose, glycerol, ethanol, and the like, or combinations thereof. Preferably, the VLA4 binding agent, if an antibody or antibody derivative, will be administered at a dose ranging between about 0.1 mg/kg body weight/day and about 20 mg/kg body weight/day, preferably ranging between about 0.1 mg/kg body weight/day and about 10 mg/kg body weight/day and at intervals of every 1-14 days. For non-antibody or antibody derivative binding agents, the dose range should preferably be between molar equivalent amounts to these amounts of antibody. Preferably, an antibody composition is administered in an amount effective to provide a plasma level of antibody of at least 1 μ g/ml. Optimization of dosages can be determined by

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administration of the binding agents, followed by
assessment of the coating of VLA4-positive cells by the
agent over time after administered at a given dose in
vivo. Peripheral blood mononuclear cells contained in
5 a sample of the individual's peripheral blood should be
probed for the presence of the agent in vitro (or ex
vivo) using a second reagent to detect the administered
agent. For example, this may be a fluorochrome
labelled antibody specific for the administered agent
10 which is then measured by standard FACS (fluorescence
activated cell sorter) analysis. Alternatively,
presence of the administered agent may be detected in
vitro (or ex vivo) by the inability or decreased
ability of the individual's cells to bind the same
15 agent which has been itself labelled (e.g., by a
fluorochrome). The preferred dosage should produce
detectable coating of the vast majority of VLA4-
positive cells. Preferably, coating is sustained in
the case of a monoclonal antibody or monoclonal
20 antibody derivative for a 1-14 day period.

In practicing this invention, treatment with VLA4
binding agents is preferably continued for as long as
the prediabetic subject maintains a stable
normoglycemic plasma level and a stable prediabetic
25 state as reflected by a number of known markers as
described above. In the Examples which follow, it has
been found that anti-VLA4 mAb, e.g., R1-2 mAb,
administration prevented diabetes onset during
treatment and that the residual beneficial results of
30 treatment were extended as long as two months following
cessation of R1-2 treatment. To sustain the full
protective effect of the VLA4 binding agent against
diabetes onset, however, continuous treatment with the
binding agents is preferred.

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Th method of the pres nt inventi n comprises administering to a pr diabetic individual a composition comprising an anti-VLA4 antibody. The examples below set forth the results observed in a rodent model of disease. These results demonstrate a protective effect of anti-VLA4 antibody in disease onset in the acute transfer model of the disease. The non-obese diabetic (NOD) mouse has become an important model of type I or insulin dependent diabetes mellitus since its introduction by Makino et al., 1980 [7] and has been documented as a particularly relevant model for human diabetes (see, e.g., Castano and Eisenbarth [1], Miller et al., 1988 [5], Hutchings et al., 1990 [33] and references cited therein). That the diabetic syndromes displayed in the NOD mouse and human are similar has been shown by several lines of evidence. For example, in both the NOD mouse and human [1], there is a strong genetic association of diabetes with loci of the major histocompatibility complex. In addition, for example, in both species, an autoimmune pathogenesis is evidenced by (i) the presence of lymphocytic inflammation in the pancreatic islets (i.e., insulitis) that appears to mediate the selective destruction of β cells, (ii) the presence of anti-islet cell antibodies, and (iii) the modulating effects of cyclosporin A. Further evidence in the NOD mouse for an autoimmune etiology of diabetes disease is (i) the ability to transfer diabetes with spleen cells (including purified splenic T cells) from diabetic donors, (ii) prevention of diabetes by *in vivo* treatment with antibodies specific for T cells, and (iii) failure of a thymic nude mice with NOD genetic background to develop moulitis or diabetes (see, e.g., Miller et al., 1988 [5], Hutchings et al., 1990 [33] and references cited therein).

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Although the precise events resulting in diabetes remain unclear, in the NOD mouse a progressive inflammatory response in the pancreas appears to be the initial histological lesion which begins as a periductal /perivascular mononuclear cell infiltrate at 3-4 weeks of age. At about 4-6 weeks of age, insulinitis may be observed and beginning at about 12 weeks of age, overt diabetes (i.e., consistent values of 1+ or higher using a Testape (Eli Lilly, Indianapolis, IN) assay for glycosuria or greater than 250 mg/dL if plasma glucose is monitored) occurs. To avoid variations in the immune status of the animals, the NOD mice are obtained from a specific pathogen-free colony and exhibit stable, high incidence of diabetes of about 80% of females and 20% of males which typically become diabetic by about 20 weeks of age. The preferred source for the NOD mice used in the experiments described herein is Taconic Farms (Germantown, NY). A large body of data, particularly from studies of the BB rat and NOD mouse has indicated that type I diabetes may be a T-cell mediated disease. Evidence to date suggests an important role for both major T cell subpopulations (CD4/L3T4 and CD8/Ly2) in the development of diabetes in man and in the NOD mouse. The data supporting the essential role of T cells in diabetes do not exclude the possibility that T lymphocytes may recruit other cells (e.g., macrophages) as the final effectors for β cell destruction. Macrophages have been implicated in the disease process based on their presence in the infiltrated islet and the ability of chronic silica treatment to prevent disease (see, e.g., Hutchings et al., 1990 [33] and references cited therein).

Using the NOD strain of mice, investigators have developed an acute transfer model of disease which

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parallels the spontaneous disease model in that transferred cells derived from diabetic NOD mice mediate the disease process, which is characterized by immune reactive cells that mediate insulinitis and islet β cell-specific destruction. Moreover, in this model, certain monoclonal antibodies against T cells (see, e.g., Miller et al., 1988 [5]) and macrophages (see, e.g., Hutchings et al., 1990 [33]) have been shown to abrogate disease onset. Such monoclonal antibodies have been used in the treatment of spontaneous disease and adoptively transferred disease, for example, anti-CD4 antibody has been shown to abrogate disease in both models (Miller et al., 1988 [5] and Shizuru et al., 1988 [30]). Results of treatment with an agent in the adoptive transfer model or spontaneous disease model are indicative of the ability of the agent to modulate the human disease process.

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EXAMPLE 1Effect of Anti-VLA4 Antibody Treatment
on Adoptive Transfer of Diabetes

For the adoptive transfer of diabetes experiments,
5 NOD mice were obtained from Taconic Farms (Germantown,
NY) or from the Joslin Diabetes Center (Boston, MA).
Spontaneously diabetic (D) females of recent onset (13-
20 weeks of age) were used as spleen cell donors and 8
week old nondiabetic (Y) females served as recipients.
10 Spleen cells from 4 week old nondiabetic (Y) female
donors which fail to transfer disease were used as a
negative control.

Recipient mice were placed on acidified water
(1:8400 dilution of concentrated HCl in water) one week
15 prior to sublethal irradiation (775 rad) performed in
split doses (300 rad, 300 rad, and 175 rad) on each of
three days (day -2, -1, and the day of transfer), in
order to minimize any incidence of intestinal infection
subsequent to high dose irradiation (Gamma Cell 1000
20 Cesium ¹³⁷ source, Nordion International, Inc., Ontario,
Canada). Spleens were harvested from diabetic donors
or from nondiabetic controls, cell suspensions made and
red cells lysed with Hemolytic Geys solution. Spleen
cells were injected intravenously ($2-3 \times 10^7$ in 0.2 ml
25 PBS) pretreated with either 75 μ g R1-2 monoclonal
antibody (mAb), 75 μ g rat IgG2b, or untreated. For the
antibody treatment, cells were simply suspended at $1-1.5 \times 10^8$
cells/ml with mAb at 375 μ g/ml and kept on
ice until injection. The timing of injection was
30 within 3 hours after last irradiation. Some recipients
received PBS alone. The anti-VLA4 mAb R1-2 and
isotype-matched rat IgG2b were purchased from
Pharmingen (La Jolla, CA). The R1-2 (rat anti-mouse)
anti-VLA4 mAb was originally described by Holzmann et

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al., 1989 [53]. The R1-2 anti-VLA4 mAb blocks VLA4 binding to its ligands (Hession et al., 1992 [54]) and therefore belongs by definition to the B group (Pulido et al., 1991 [44], i.e., is equivalent to anti-human VLA4 mAbs of the B group (e.g., HP1/2 or HP2/1).

The R1-2 mAb or rat IgG2b was administered at a dose of 75 μ g/0.2 ml intraperitoneally every 2-3 days, a dosing regimen which was determined to maintain maximal coating of VLA4-positive cells in the peripheral blood, lymphoid organs and bone marrow as detected by staining of peripheral blood cells and single cell suspensions prepared from these organs with a fluorochrome labelled mAb specific for the R1-2 mAb and FACS analysis to measure fluorochrome positive cells (as described above). Injections were maintained through day 12 or day 24 post transfer. Mice were monitored for diabetes by testing for glycosuria with TestTape (Eli Lilly, Indianapolis, IN) and by plasma glucose levels (Glucometer, 3 Blood Glucose Meter, Miles, Inc., Elkhart, IN) and were considered diabetic after two consecutive urine positive tests [TestTape values of [+1] or higher] or plasma glucose levels >250 mg/dL.

An inhibitory effect of the anti-VLA4 mAb on the onset of diabetes was demonstrated when spleen cells isolated from NOD diabetic donors were treated with a saturating quantity of anti-VLA4 mAb R1-2 followed by transfer into nondiabetic irradiated hosts, as described above, and the R1-2 mAb was then administered every other day for 12 days in order to maintain maximal coating of all VLA4-positive cells in the peripheral blood and lymphoid organs for two weeks. Figure 1 shows the frequency of recipients that became diabetic and the day of disease onset for transfer of 2×10^7 splenocytes from diabetic NOD donor (D \rightarrow Y) (i)

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without treatment (closed circles); (ii) with rat IgG2b
treatment (closed triangles), and (iii) with R1-2 anti-
VLA4 treatment (closed diamonds) as well as for
transfer of splenocytes from non-diabetic NOD donors
(Y→Y) (open squares). Injection of PBS alone gave 0%
incidence. Under these conditions, only 1 of 8
individual R1-2 mAb treated recipients became diabetic,
with onset on day 29 post transfer. By contrast, 6/10
and 5/9 individuals became diabetic after receiving
splenocytes from diabetic donors treated with no mAb or
with non-specific rat IgG2b, respectively. As shown in
Figure 1, diabetes onset occurred as early as day 14
post transfer, though administration of the irrelevant
rat IgG2b somewhat delayed onset.

These data demonstrate a protective effect of the
R1-2 mAb which was dependent upon its specificity for
VLA4. Recipients of splenocytes from nondiabetic mice
or of PBS alone failed to become diabetic. Thus,
treatment with anti-VLA4 antibody reduced the frequency
of diabetes during 30 days post transfer.

Although the results shown in Figure 1 demonstrate
that clinical diabetes occurred in only 1 of 8 anti-
VLA4 treated animals, it was possible that the anti-
VLA4 antibody caused only a minor delay in the onset of
disease. Plasma glucose levels were monitored in
parallel with urine glucose in order to quantify any
increase in blood sugar levels and thereby detect
progression to clinical disease. In the anti-VLA4
antibody treated group shown in Figure 1, all mice were
still normoglycemic on day 29 with an average plasma
glucose value of 100 ± 7 mg/dL, $n=7$, except for the
single individual who scored as clinically diabetic by
urine test and plasma glucose >500 mg/dL. Thus,
disease progression was not apparent in any of the
other anti-VLA4 antibody treated recipients shown in

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Figure 1 on day 29 post transfer, a full 2 weeks beyond the last anti-VLA4 antibody injection. Analysis of sera from these mice confirmed that the anti-VLA4 mAb dropped to low or undetectable levels by day 18-21 post-transfer.

Additional cell transfers were performed in order to confirm that the anti-VLA4 mAb protected against transfer of diabetes. In these experiments, the anti-VLA4 antibody treatment was extended to day 25 post transfer but administered every 3.5 days thereby maintaining saturating levels of R1-2 mAb or rat IgG2b through day 26 when mice were sacrificed for pancreatic tissue. Under these conditions, an inhibitory effect of the anti-VLA4 mAb on the onset of diabetes was also demonstrated upon spleen cell transfer and R1-2 treatment. Figure 2 shows the frequency of recipients (n=4-5 for each group) that became diabetic and the day of disease onset for transfer of 3×10^7 splenocytes from diabetic NOD donors (D→Y) (i) without treatment (closed circles), (ii) with IgG2b treatment (closed triangles) and with R1-2 anti-VLA4 treatment (closed diamonds), as well as for transfer of splenocytes from non-diabetic NOD donors (Y→Y; open squares). Injection of PBS alone gave 0% incidence. Figure 2 shows that only 1 out of 5 R1-2 mAb treated mice became diabetic by day 22 post transfer whereas diabetes was transferred in 4/4 recipients without R1-2 mAb and 5/5 treated with rat IgG2b. Disease onset occurred as early as day 13 post transfer. These experiments, individually and collectively demonstrate that anti-VLA4 mAb reproducibly protects against development of diabetes in an acute transfer model of disease.

Further experiments were performed to determine whether the anti-VLA-4 mAb simply delayed disease onset during the treatment period or if it could achieve a

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longer-term protective effect. Figure 3 shows the onset of diabetes in mice over time after R1-2 injection (once every 3.5 days through day 25) with only 2/5 mice becoming diabetic on days 35 and 38 post transfer, 10-13 days after the last R1-2 injection. By contrast, diabetes occurred in the untreated and IgG2b treated groups as early as day 11 post transfer, with 100% incidence by days 18-21. Surprisingly, disease incidence in the R1-2 treated group did not further increase even as long as 2 months following the last R1-2 injection. Plasma glucose values monitored in parallel during this time reveal that these three individuals were consistently normoglycemic. After this point (i.e., approximately 3 months post-transfer), even the negative control groups which received PBS alone or non-diabetic cells begin developing spontaneous disease. In summary, the VLA-4-specific mAb reduces the incidence of diabetes transfer. Moreover, its protective effect against disease is sustained in the absence of further mAb treatment.

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EXAMPLE 2Eff ct of Anti-VLA4 mAb on Pancreatis Insulitis

For histological analysis, mice were sacrificed between 2-4 weeks post-transfer as described in this Example and pancreata harvested in 10% formalin buffered saline for paraffin-embedded sections which were stained with hematoxylin and eosin (H&E) for histology. Degree of insulitis was scored as follows: Grade 0: no insulitis [islet devoid of inflammation]; Grade I: peri-insulitis [inflammatory mononuclear cells located peripheral to the islet]; Grade II: <25% infiltrated [<25% of the islet interior contains lymphocytic inflammatory cells]; Grade III: 25-50% infiltrated [lymphocytic infiltration]; Grade IV: >50% infiltrated. The percent of islets in each Grade was then calculated relative to the total number of islets examined. Histologic sections were examined and scored for the degree of insulitis following the adoptive transfer of NOD splenocytes with and without anti-VLA4 mAb treatment and the results tabulated. Specifically, the frequency of uninfiltrated islets (Grade 0-I infiltrate) and islets with Grade II-IV insulitis (as described above) were quantitated. For each experimental group, pancreatic sections from n= 4-5 mice were scored.

Pancreatic tissue was recovered from recipients treated with the anti-VLA-4 mAb for various time periods in order to address its effect on the establishment of islet-specific cellular infiltrates. Mice were treated with nonspecific rat IgG2b or R1-2 mAb every 3.5 days through day 14 when sacrificed. Similarly, mice were treated through day 25 and sacrificed after diabetes was diagnosed or on day 26 post transfer. Mice continuously treated with the R1-2 mAb for 14 days post transfer maintain a high frequency

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(76%) of uninfilitrated islets, with only 24% progressing to grade II-IV insulinitis. By contrast those treated with nonspecific rat IgG2b show the reciprocal pattern, with 74% severe insulinitis.

5 Likewise, in the mice treated with R1-2 through day 25 (20% diabetic, pancreata isolated from mice reported in Figure 2), a high frequency (58%) of uninfilitrated islets were preserved, similar to that (55% uninfilitrated) in nondiabetic recipients of young NOD
10 splenocytes, as shown in Figure 4. By contrast, both the untreated or IgG2b-treated mice had only 28% uninfilitrated islets, and conversely had increased (72%) insulinitis. Thus, the anti-VLA-4 mAb treatment appears to specifically inhibit or alternatively to
15 delay the development of insulinitis upon adoptive transfer of diabetogenic spleen cells.

In order to distinguish between these alternatives, the pattern of insulinitis after 4 weeks post transfer was determined when mice were treated
20 with rat IgG2b or R1-2 mAb through day 12 and then maintained without further treatment. Mice were sacrificed upon diabetes diagnosis or on day 29 post transfer. Analysis of sera from these mice confirmed that circulating anti-VLA-4 mAb dropped to undetectable
25 levels by days 18-21 post transfer. With this protocol, the degree of insulinitis in the R1-2-treated group (69% insulinitis, 25% diabetic) was similar to that in untreated recipients (73% insulinitis, 60% diabetic) though still lower than that in the rat IgG2b-treated
30 mice (96% insulinitis, 75% diabetic), as shown in Figure 5. Significantly, the severity of insulinitis was similar between the R1-2 treated, untreated and rat IgG2b treated groups with an average of 57%, 47%, 64% Grade III/IV infiltrates, respectively. Even
35 considering only the nondiabetic R1-2 treated

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individuals, they still exhibited 59% insulinitis with 52% Grade III/IV infiltrates. Recipients of nondiabetogenic NOD splenocytes had only 7% Grade III/IV infiltrates. Conversely, Figure 5 shows that the frequency of uninfilitrated islets was decreased in the R1-2 treated mice as compared to recipients of saline or nondiabetogenic spleen cells. Thus, the degree of insulitis progressed in these R1-2 treated mice (Figure 5) as compared to mice wherein R1-2 treatment was maintained (Figure 4) and approached that in the untreated and rat IgG2b treated control groups. Taken together, these data indicate that anti-VLA-4 mAb administration can delay the progression of insulitis in an acute transfer model of disease.

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EXAMPLE 3**Comparison of Different Anti-VLA4 Antibody
Treatment in Adoptive Transfer of Diabetes**

5 This Example provides comparative efficacy results
of PS/2, an anti-VLA4 antibody, with R1-2 using the
adoptive transfer model and procedure described in
Example 1. NOD mice were treated with (a) an
irrelevant control antibody (D/rat IgG2b, n = 19 mice);
10 (b) R1-2 antibody (D/R1-2 mAb, n = 24 mice); (c) PS/2
mAb (D/PS/2 mAb, n = 5 mice); or (d) no treatment
(NONE, n = 26 mice). Spleen cells were injected
intravenously ($2-3 \times 10^7$ in 0.2 ml PBS) and pretreated
with either 75 μ g R1-2 mAb, 75 μ g PS/2 mAb, 75 μ g rat
IgG2b, or untreated. Isolation and purification of
15 PS/2 anti-VLA4 mAb was originally described by Miyake
et al., 1991 [55].

The R1-2 mAb, PS/2 mAb or rat IgG2b was
administered at a dose of 75 μ g/0.2 ml
intraperitoneally every 2-3 days, a dosing regimen
20 which was determined to maintain maximal coating of
VLA4-positive cells in the peripheral blood, lymphoid
organs and bone marrow as detected by staining of
peripheral blood cells and single cell suspensions
prepared from these organs with a fluorochrome labelled
25 mAb specific for the R1-2 and PS/2 mAb and FACS
analysis to measure fluorochrome positive cells (as
described above). Injections were maintained through
days 22 to 25 post transfer. Mice were monitored for
diabetes by testing for glycosuria with TestTape (Eli
30 Lilly, Indianapolis, IN) and by plasma glucose levels
(Glucometer, 3 Blood Glucose Meter, Miles, Inc.,
Elkhart, IN) and were considered diabetic after two
consecutive urine positive tests [Testape values of
[+1] or higher] or plasma glucose levels >250 mg/dL.

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An inhibitory effect of the anti-VLA4 mAb on the onset of diabetes was demonstrated when splenic cells isolated from NOD diabetic donors were treated with a saturating quantity of anti-VLA4 mAb R1-2 or PS/2 followed by transfer into nondiabetic irradiated hosts, as described above, and the R1-2 mAb or PS/2 mAb was then administered every other day for 22-25 days in order to maintain maximal coating of all VLA4-positive cells in the peripheral blood and lymphoid organs for about two weeks. Table 1 shows the frequency of recipients that became diabetic and the day of disease onset for transfer of splenocytes from diabetic NOD donor (i) without treatment (D); (ii) with rat IgG2b treatment (D/nonspecific rat IgG2b); (iii) with R1-2 anti-VLA4 treatment (D/R1-2 mAb); (iv) with PS/2 treatment (D/PS/2 mAb) as well as for transfer of splenocytes from non-diabetic NOD donors (non-D). Non-diabetic mice receiving PBS and no splenocytes (NONE) were included as a control. Injection of PBS alone gave 4% incidence. Under these conditions, only 1 of 24 individual R1-2 mAb treated recipients became diabetic, with onset on day 22 post transfer while none of the five individual PS/2 mAb treated recipients became diabetic. By contrast, 16/19 individuals became diabetic after receiving splenocytes from diabetic donors treated with no mAb or with non-specific rat IgG2b. As shown in Table 1, diabetes onset occurred as early as day 14 post transfer, though administration of the irrelevant rat IgG2b somewhat delayed onset by one day.

These data demonstrate a protective effect of the R1-2 mAb and PS/2 which were dependent upon its specificity for VLA4. Recipients of splenocytes from nondiabetic mice or of PBS alone failed to become diabetic. Thus, treatment with anti-VLA4 antibody

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reduced the frequency of diabetes during 30 days post transfer. Analysis of sera from these mice confirmed that levels of R1-2 and PS/2 anti-VLA4 mAb become undetectable between days 26 and 34 post-transfer.

5

TABLE 1

**Anti-VLA-4 mAbs Inhibit Adoptive
Transfer of Diabetes in NOD Mice**

	Cells Transferred/Treatment*	No. Diabetic/Total Recipients*		Day of Onset X \pm SEM
10	NONE	1/26	(4%)	34
	Non-D	1/15	(7%)	15
	D	16/19	(84%)	14 \pm 0.2
	D/Nonspecific rat IgG2b	16/19	(84%)	15 \pm 0.9
15	D/R1-2 mAb	1/24	(4%)	22
	D/PS/2 mAb	0/5	(0%)	

*Spleen cells from 4 week old nondiabetic (NON-D) or from new onset diabetic (D) NOD females were transferred, with D cells suspended in mAb or rat IgG or without mAb before transfer and recipients treated twice weekly for 22-25 days. Mice were monitored for one month post transfer. Data are compiled from 5 experiments.

*D/R1-2 and D/PS/2 mAb treated groups are significantly different from D and D/rat IgG2b treated groups by Chi square test with Yates' correction as follows: R1-2 vs. IgG2b treated and D group, $p < 0.0001$; PS/2 vs. IgG2b treated and D group, $p < 0.003$.

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EXAMPLE 4**Effect of Anti-VLA4 Antibody Treatment
in Spontaneous Diabetes Model**

5 This Example described efficacy results using R1-2
mAb in the spontaneous diabetes model which employs NOD
mice. NOD mice were treated for 8 weeks with (a) an
irrelevant control antibody (NOD/rat IgG2b, n = 10
mice); (b) R1-2 antibody (NOD/R1-2, n = 20 mice); or
10 (c) no treatment (NOD, n = 10 mice) starting at week
four to week twelve of age. mAb was administered at a
dose of 75 μ g in 0.2 ml PBS iv, twice weekly. Mice
were monitored for diabetic events by TestTape for
glycosuria as previously described.

15 Figure 6 demonstrates a marked delay in diabetes
onset (12-16 weeks delay) following R1-2
administration, as compared to the two control groups.
NOD mice which received irrelevant IgG2b mAb or no
treatment developed diabetes as early as 13 weeks.
These spontaneous disease model results parallel the
20 adoptive transfer results with R1-2 mAb illustrated in
Figure 1 and directly demonstrate that an anti-VLA4
antibody protects against diabetes onset.

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EXAMPLE 5**Effect of a VCAM-Ig Fusion Protein
on Adoptive Transfer of Diabetes**

5 The adoptive transfer experiment described in
Example 1 was repeated with a VCAM-Ig fusion protein
(VCAM 2D-IgG) instead of an anti-VLA4 mAb. VCAM 2D-IgG
is a soluble form of the ligand for VLA4 (VCAM1) which
consists of the two N-terminal domains of VCAM1 fused
10 to the human IgG1 heavy chain constant region sequences
(Hinges, C_H2 and C_H3). The VCAM 2D-IgG DNA sequence
and its translated amino acid sequence are shown in SEQ
ID NO: 9. Figure 8 illustrates the fusion protein
structure. The fusion protein was constructed by
recombinant techniques as described below.

15 **Isolation of cDNA of Human IgG1 Heavy Chain
Region and Construction of Plasmid pSAB144**

In order to isolate a cDNA copy of the human IgG1
heavy chain region, RNA was prepared from COS7 cells
which has been transiently transfected by the plasmid
20 VCAM1-IgG1 (also known as pSAB133). Construction of
plasmid VCAM1-IgG1 is described in PCT patent
application WO 90/13300. The RNA was reverse
transcribed to generate cDNA using reverse
transcriptase and random hexamers as the primers.
25 After 30 min. at 42°C, the reverse transcriptase
reaction was terminated by incubation of the reaction
at 95°C for 5 min. The cDNA was then amplified by PCR
(Polymerase Chain Reaction, see, e.g., Sambrook et al.,
Molecular Cloning, Vol. 3, pp. 14.1-14.35 (Cold Spring
30 Harbor; 1989)) using the following kinased primers:
370-31 (SEQ ID NO: 10):

5'TCGTC GAC AAA ACT CAC ACA TGC C
Asp Lys Thr His Thr Cys

which contains a SalI site, and

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370-32 (SEQ ID NO: 11):

5' GTAAATGAGT GCGGCGGCCG CCAA,

which encodes the carboxy terminal lysine of the IgG1 heavy chain constant region, followed by a NotI site.

5 The PCR amplified cDNA was purified by agarose gel electrophoresis and glass bead elution for cloning in plasmid pNN03. Plasmid pNN03 was constructed by removing the synthetic polylinker sequence from the commercially available plasmid pUC8 (Pharmacia, Piscataway, New Jersey) by restriction endonuclease digestion and replacing the synthetic polylinker sequence with the following novel synthetic sequence (SEQ ID NO: 12):

10 GCGGCCGCGG TCCAACCACC AATCTCAAAG CTTGGTACCC GCGAATTCAG
15 ATCTGCAGCA TGCTCGAGCT CTAGATATCG ATTCCATGGA TCCTCACATC
 CCAATCCGCG GCCGC.

 The purified PCR amplified cDNA fragment was ligated to pNN03 which had been cleaved with EcoRV, dephosphorylated, and purified by low melt agarose gel electrophoresis. The ligation reaction was used to transform E.coli JA221 and the resulting colonies were screened for a plasmid containing an insert of approximately 700 bp. The identity of the correct insert was confirmed by DNA sequence analysis, and the plasmid was designated pSAB144.

Construction of Plasmid pSAB142

 The plasmid pSAB142 was constructed as follows. CDNA prepared from COS cells transfected with pSAB133 (as described in the previous section) was subjected to PCR amplification using oligonucleotides 370-01 and 370-29. Oligonucleotide 370-01 includes a NotI site and the nucleotides corresponding to amino acids 1 through 7 of the VCAM-1 signal sequence

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(SEQ ID NO: 13):

5' GAGCTCGAGGCGGCCGCACCATGCCTGGGAAGATGGTCGTG
MetProGlyLysMetValVal

5 Oligonucleotide 370-29 corresponds to the VCAM-1 amino acids 214-219 and includes a SalI site (SEQ ID NO: 14):

5'AA GTC GAC TTG CAA TTC TTT TAC

The amplified DNA fragment was ligated to the vector fragment of pNN03, cleaved by EcoRV.

Construction of pSAB132

10 pJOD-S (Barsoum, J., DNA and Cell Biol., 2, pp.293-300 (1990)) was modified to insert a unique NotI site downstream from the adenovirus major late promoter so that NotI fragments could be inserted into the expression vector. pJOD-S was linearized by NotI
15 cleavage of the plasmid DNA. The protruding 5' termini were blunt-ended using Mung Bean nuclease, and the linearized DNA fragment was purified by low melting temperature agarose gel electrophoresis. The DNA fragment was religated using T4 DNA ligase. The
20 ligated molecules were then transformed into E.coli JA221. Colonies were screened for the absence of a NotI site. The resulting vector was designated pJOD-S delta Not1. pJOD-8 delta Not1 was linearized using SalI and the 5' termini were dephosphorylated using
25 calf alkaline phosphatase. The linearized DNA fragment was purified by low melting temperature agarose gel electrophoresis and ligated in the presence of phosphorylated oligonucleotide ACE175, which has the following sequence (SEQ ID NO:15):

30 TCGACGCGGC CGCG

The ligation mixture was transformed into E.coli JA221, and colonies were screened for the presence of a

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plasmid having a NotI site. The desired plasmid was named pMDR901.

5 In order to delete the two SV40 enhancer regions in the Sv40 promoter which controls transcription of the DHFR cDNA, pMDR901 and pJODae-TPA (Barsoum, DNA and Cell Biol., 9, pp. 293-300 (1990)), both were cleaved with AatII and DraIII. The 2578 bp AatII-DraIII fragment from pMDR901 and the 5424 bp AatII-DraIII fragment from pJODae-TPA were isolated by low melting temperature agarose gel electrophoresis and ligated together. Following transformation into E.coli JA221, the resulting plasmid, pMDR902, was isolated. pSAB132 was then formed by eliminating the EcoRI-NotI fragment of pMDR902 containing the adenovirus major late promoter and replacing it with an 839 bp EcoRI-NotI fragment from plasmid pCMV-B (Clontech, Palo Alto, California) containing the human cytomegalovirus immediate early promoter and enhancer.

Construction of pSAB146

20 pSAB144 was cleaved with SalI and NotI, and the 693 bp fragment isolated. pSAB142 was cleaved with NotI and SalI and the 664 bp fragment was isolated. The two fragments were ligated to pSAB132 which had been cleaved with NotI, and the 5' termini
25 dephosphorylated by calf alkaline phosphatase. The resulting plasmid, pSAB146, contained the DNA sequence encoding the VCAM-1 signal sequence, the amino terminal 219 amino acids of mature VCAM-1, ten amino acids of the hinge region of IgG1 and the C_H2 and C_H3 constant domains of IgG1.
30

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Production of VCAM 2D-IgG from a stably transformed CHO cell line

5 A recombinant VCAM 2D-IgG expression vector was constructed as described below and transfected into CHO cells to produce a cell line continuously secreting VCAM 2D-IgG.

10 The 1.357 kb NotI fragment containing the VCAM 2D-IgG coding sequence of pSAB146 was purified by agarose gel electrophoresis. This fragment was ligated into the NotI cloning site of the expression vector pMDR901, which uses the adenovirus 2 major late promoter for heterologous gene expression and the selectable, amplifiable dihydrofolate reductase (dhfr) marker. The ligated DNA was used to transform E.coli DH5. Colonies
15 containing the plasmid with the desired, correctly oriented insert were identified by the presence of 5853 and 3734 bp fragments upon digestion with Hind III; and 4301, 2555, 2293, and 438 bp fragments upon digestion with Bgl II. The resultant recombinant VCAM 2D-IgG
20 expression vector was designated pEAG100. The identity of the correct insert was confirmed by DNA sequence analysis.

The recombinant expression plasmid pEAG100 was electroporated into dhfr-deficient CHO cells according
25 to the published protocol of J. Barsoum (DNA Cell Biol 9: 293-300, 1990), with the following changes: 200 µg of Pvu I-linearized pEAG100 plasmid and 200 µg of sonicated salmon sperm DNA were used in the electroporation protocol. In addition, cells were
30 selected in alpha-complete medium supplemented with 200 nM methotrexate.

To determine expression levels of secreted VCAM 2D-IgG, clones were transferred to a flat bottom 96 well microtiter plate, grown to confluency and assayed
35 by ELISA as described below.

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Wells of Immulon 2 plates (Dynatech, Chantilly, Virginia) were each coated with anti-VCAM MAb 4B9 (isolated and purified on Protein A Sepharose as described by Carlos et al, 1990 [56]) with 100 μ l of anti-VCAM 4B9 MAb diluted to 10 μ g/ml in 0.05 M sodium carbonate/bicarbonate buffer, pH 9.6, covered with Parafilm, and incubated overnight at 4°C. The next day, the plate contents were dumped out and blocked with 200 μ l/well of a block buffer (5% fetal calf serum in 1X PBS), which had been filtered through a 2 μ filter. The buffer was removed after a 1 hour incubation at room temperature and the plates were washed twice with a solution of 0.05% Tween-20 in 1X PBS. Conditioned medium was added at various dilutions. As a positive control, an anti-mouse Ig was also included. Block buffer and LFA-3TIP constituted as negative controls. The samples and controls were incubated at room temperature for 2 hours.

The plates were then washed twice with a solution of 0.05% Tween-20 in 1X PBS. Each well, except for the positive control well, was then filled with 50 μ l of a 1:2000 dilution of HRP-Donkey anti-human IgG (H+L) (Jackson Immune Research Laboratories, Inc.; West Grove, Pennsylvania) in block buffer. The positive control well was filled with 50 μ l of a 1:2000 dilution of HRP-Goat anti-mouse IgG (H+L) (Jackson Immune Research Laboratories, Inc.; West Grove, Pennsylvania) in block buffer. The plates were then incubated for 1 hour at room temperature.

The HRP conjugated Ab solutions were removed, and the wells were washed twice with 0.05% Tween-20 in 1X PBS. Then, 100 μ l of HRP-substrate buffer was added to each well at room temperature. HRP-substrate buffer was prepared as follows: 0.5 ml of 42mM 3,3', 5,5'-tetramethylbenzidine (TMB), (ICN Immunobiologicals,

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5 Lisle, South Carolina, Catalogue No. 980501) in DMSO (Aldrich) was slowly added to 50 ml of substrate buffer (0.1 M sodium acetate/citric acid, pH4.9); followed by addition of 7.5 μ l of 30% hydrogen peroxide (Sigma, Catalogue No. H-1009).

10 The development of a blue color in each well was monitored at 650nm on a microtiter plate reader. After 7-10 minutes, the development was stopped by the addition of 100 μ l of 2N Sulfuric acid. The resulting yellow color was read at 450nm on a microtiter plate reader. A negative control well was used to blank the machine.

Purification of VCAM 2D-IgG

15 CHO cells expressing VCAM 2D-IgG were grown in roller bottles on collagen beads. Conditioned medium (5 Liters) was concentrated to 500 ml using an Amicon S1Y10 spiral ultrafiltration cartridge (Amicon, Danvers, MA). The concentrate was diluted with 1 liter of Pierce Protein A binding buffer (Pierce, Rockford, IL) and gravity loaded onto a 10 ml Protein A column (Sephacrose 4 Fast Flow, Pharmacia, Piscataway, NJ).
20 The column was washed 9 times with 10 ml of Protein A binding buffer and then 7 times with 10 ml of PBS. VCAM 2D-IgG was eluted with twelve 5 ml steps
25 containing 25 mM H_3PO_4 , pH2.8, 100 mM NaCl. The eluted samples were neutralized by adding 0.5 M Na_2HPO_4 , pH8.6 to 25 mM. Fractions were analyzed for absorbance at 280 nm and by SDS-PAGE. The three peaks fractions of highest purity were pooled, filtered, aliquoted and
30 stored at -70°C. By SDS-PAGE, the product was greater than 95% pure. The material contained less than 1 endotoxin unit per mg of protein. In some instances, it was necessary to further purify the Protein A eluate product on Q-Sepharose FF (Pharmacia). The protein A

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uate was diluted with 3 volumes of 25 mM Tris HCl pH 8.0 and loaded onto a Q-Sepharose FF column at 10 mg VCAM 2D-IgG per ml for 15 min. The VCAM 2D-IgG was then eluted from the Q-Sepharose with PBS.

5 Evaluation of VCAM 2D-IgG

 Spleen cell suspensions were prepared from diabetic donors or from nondiabetic controls as described above. Spleen cells were injected intravenously ($2-3 \times 10^7$ in 0.2 ml PBS) and were pretreated with either 100 μ g VCAM 2D-IgG or 100 μ g of irrelevant LFA-3Ig fusion protein control. Another group received PBS alone without cells transferred. The fusion protein LFA-3Ig (LFA-3TIP) was isolated and purified as described in PCT US92/02050 and Miller et al., 1993 [57]. The VCAM 2D-IgG fusion protein or irrelevant LFA-3Ig protein was administered at a dose of 100 μ g/0.2 ml intraperitoneally twice weekly through day 17. This concentration was sufficient to provide a serum level of fusion protein sufficient to saturate VLA4-positive cells, the serum levels determined by ELISA as described above. Diabetes onset was monitored as described above.

 The results of the evaluation are shown in Figure 7. As shown in this Figure, VCAM 2D-IgG fusion protein significantly inhibits the onset of diabetes in recipients of cells from diabetic donor mice (D/VCAM-Ig, open circles) with 60% incidence by day 30 post-transfer, as compared to the mice which received cells from diabetic donor (data not shown) and LFA-3Ig irrelevant control Ig fusion protein (D/LFA-3 Ig) which had already achieved 60% incidence by day 15 post-transfer. Mice which received no cells (PBS only) did not develop disease. There were $n = 5$ mice per experimental group.

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In summary, VLA4 binding agents such as anti-VLA4 antibodies were protective against diabetes disease onset (Examples 1, 3 and 4) and were effective in delaying the progression of insulinitis (Example 2) using a murine model for human diabetes. Other VLA4 binding agents such as soluble VCAM derivatives (VCAM 2D-IgG) were also useful in protecting against diabetes disease onset (Example 5). The foregoing examples are intended as an illustration of the method of the present invention and are not presented as a limitation of the invention as claimed hereinafter. From the foregoing disclosure, numerous modifications and additional embodiments of the invention will be apparent to those experienced in this art. For example, actual dosage used, the type of antibody or antibody fragment used, mode of administration, exact composition, time and manner of administration of the treatment, and many other features all may be varied without departing from the description above. All such modifications and additional embodiments are within the contemplation of this application and within the scope of the appended claims.

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- 50 -

The foregoing documents are incorporated herein by reference in their entirety.

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SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Burkly, Linda C.
- (ii) TITLE OF INVENTION: Treatment for Insulin Dependent Diabetes
- (iii) NUMBER OF SEQUENCES: 15
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Allegretti & Witcoff, Ltd.
 - (B) STREET: 10 South Wacker Drive, Suite 3000
 - (C) CITY: Chicago
 - (D) STATE: IL
 - (E) COUNTRY: US
 - (F) ZIP: 60606
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: Word Perfect 5.1
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/029,330
 - (B) FILING DATE: 9 February 1993
 - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Miao, Emily
 - (B) REGISTRATION NUMBER: 35,285
 - (C) REFERENCE/DOCKET NUMBER: 92,749-A; D015 CIP PCT
- (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 312-715-1000
 - (B) TELEFAX: 312-715-1234

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 360 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (ix) FEATURE:

- 52 -

- (A) NAME/KEY: misc_feature
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 but Gln (Q) may be substituted"

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- (A) NAME/KEY: CDS
 (B) LOCATION: 1-360

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TCA	GTC	AAG	TTG	TCC	TGC	ACA	GCT	TCT	GGC	TTC	AAC	ATT	AAA	GAC	90
Ser	Val	Lys	Leu	Ser	Cys	Thr	Ala	Ser	Gly	Phe	Asn	Ile	Lys	Asp	
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ACC	TAT	ATG	CAC	TGG	GTG	AAG	CAG	AGG	CCT	GAA	CAG	GGC	CTG	GAG	135
Thr	Tyr	Met	His	Trp	Val	Lys	Gln	Arg	Pro	Glu	Gln	Gly	Leu	Glu	
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TGG	ATT	GGA	AGG	ATT	GAT	CCT	GCG	AGT	GGC	GAT	ACT	AAA	TAT	GAC	180
Trp	Ile	Gly	Arg	Ile	Asp	Pro	Ala	Ser	Gly	Asp	Thr	Lys	Tyr	Asp	
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CCG	AAG	TTC	CAG	GTC	AAG	GCC	ACT	ATT	ACA	GCG	GAC	ACG	TCC	TCC	225
Pro	Lys	Phe	Gln	Val	Lys	Ala	Thr	Ile	Thr	Ala	Asp	Thr	Ser	Ser	
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AAC	ACA	GCC	TGG	CTG	CAG	CTC	AGC	AGC	CTG	ACA	TCT	GAG	GAC	ACT	270
Asn	Thr	Ala	Trp	Leu	Gln	Leu	Ser	Ser	Leu	Thr	Ser	Glu	Asp	Thr	
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GCC	GTC	TAC	TAC	TGT	GCA	GAC	GGA	ATG	TGG	GTA	TCA	ACG	GGA	TAT	315
Ala	Val	Tyr	Tyr	Cys	Ala	Asp	Gly	Met	Trp	Val	Ser	Thr	Gly	Tyr	
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GCT	CTG	GAC	TTC	TGG	GGC	CAA	GGG	ACC	ACG	GTC	ACC	GTC	TCC	TCA	360
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- (A) LENGTH: 120 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

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(ii) MOLECULE TYPE: protein

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				36					41					46	
Trp	Ile	Gly	Arg	Ile	Asp	Pro	Ala	Ser	Gly	Asp	Thr	Lys	Tyr	Asp	
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Pro	Lys	Phe	Gln	Val	Lys	Ala	Thr	Ile	Thr	Ala	Asp	Thr	Ser	Ser	
				66					71					76	
Asn	Thr	Ala	Trp	Leu	Gln	Leu	Ser	Ser	Leu	Thr	Ser	Glu	Asp	Thr	
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Ala	Val	Tyr	Tyr	Cys	Ala	Asp	Gly	Met	Trp	Val	Ser	Thr	Gly	Tyr	
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(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 318 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1-318
- (D) OTHER INFORMATION: /product= "HP1/2 light chain variable region"

(ix) FEATURE:

- (A) NAME/KEY: misc_feature
- (B) LOCATION: 1
- (D) OTHER INFORMATION: /note= "pBAG172 insert: HP1/2 light chain variable region"

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Gly	Asp	Arg	Val	Thr	Ile	Thr	Cys	Lys	Ala	Ser	Gln	Ser	Val	Thr	
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AAT	GAT	GTA	GCT	TGG	TAC	CAA	CAG	AAG	CCA	GGG	CAG	TCT	CCT	AAA	135
Asn	Asp	Val	Ala	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	Gln	Ser	Pro	Lys	
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CTG	CTG	ATA	TAT	TAT	GCA	TCC	AAT	CGC	TAC	ACT	GGA	GTC	CCT	GAT	180
Leu	Leu	Ile	Tyr	Tyr	Ala	Ser	Asn	Arg	Tyr	Thr	Gly	Val	Pro	Asp	
				50					55					60	
CGC	TTC	ACT	GGC	AGT	GGA	TAT	GGG	ACG	GAT	TTC	ACT	TTC	ACC	ATC	225
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				65					70					75	
AGC	ACT	GTG	CAG	GCT	GAA	GAC	CTG	GCA	GTT	TAT	TTC	TGT	CAG	CAG	270
Ser	Thr	Val	Gln	Ala	Glu	Asp	Leu	Ala	Val	Tyr	Phe	Cys	Gln	Gln	
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GAT	TAT	AGC	TCT	CCG	TAC	ACG	TTC	GGA	GGG	GGG	ACC	AAG	CTG	GAG	315
Asp	Tyr	Ser	Ser	Pro	Tyr	Thr	Phe	Gly	Gly	Gly	Thr	Lys	Leu	Glu	
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ATC															
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 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

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Leu	Leu	Ile	Tyr	Tyr	Ala	Ser	Asn	Arg	Tyr	Thr	Gly	Val	Pro	Asp

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	50		55		60
Arg Phe Thr Gly Ser Gly Tyr Gly Thr Asp Phe Thr Phe Thr Ile					
	65		70		75
Ser Thr Val Gln Ala Glu Asp Leu Ala Val Tyr Phe Cys Gln Gln					
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Asp Tyr Ser Ser Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu					
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(2) INFORMATION FOR SEQ ID NO:5:

- (i) SEQUENCE CHARACTERISTICS:
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 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (ix) FEATURE:
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 - (B) LOCATION: 1-57
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 - (D) OTHER INFORMATION: /note= "pBAG195 insert: AS heavy chain variable region"
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Gly Ala His Ser Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu	
	1 5 10

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GTC AGA CCT AGC CAG ACC CTG AGC CTG ACC TGC ACC GCG TCT GGC 135
Val Arg Pro Ser Gln Thr Leu Ser Leu Thr Cys Thr Ala S r Gly
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TTC AAC ATT AAA GAC ACC TAT ATG CAC TGG GTG AGA CAG CCA CCT 180
Phe Asn Ile Lys Asp Thr Tyr Met His Trp Val Arg Gln Pro Pro
              30                      35                      40

GGA CGA GGT CTT GAG TGG ATT GGA AGG ATT GAT CCT GCG AGT GGC 225
Gly Arg Gly Leu Glu Trp Ile Gly Arg Ile Asp Pro Ala Ser Gly
              45                      50                      55

GAT ACT AAA TAT GAC CCG AAG TTC CAG GTC AGA GTG ACA ATG CTG 270
Asp Thr Lys Tyr Asp Pro Lys Phe Gln Val Arg Val Thr Met Leu
              60                      65                      70

GTA GAC ACC AGC AGC AAC CAG TTC AGC CTG AGA CTC AGC AGC GTG 315
Val Asp Thr Ser Ser Asn Gln Phe Ser Leu Arg Leu Ser Ser Val
              75                      80                      85

ACA GCC GCC GAC ACC GCG GTC TAT TAT TGT GCA GAC GGA ATG TGG 360
Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala Asp Gly Met Trp
              90                      95                      100

GTA TCA ACG GGA TAT GCT CTG GAC TTC TGG GGC CAA GGG ACC ACG 405
Val Ser Thr Gly Tyr Ala Leu Asp Phe Trp Gly Gln Gly Thr Thr
              105                      110                      115

GTC ACC GTC TCC TCA GGT GAG TCC
Val Thr Val Ser Ser Gly Glu Ser
              120

```

429

(2) INFORMATION FOR SEQ ID NO:6:

- (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 143 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

```

Met Asp Trp Thr Trp Arg Val Phe Cys Leu Leu Ala Val Ala Pro
              -15                      -10                      -5

Gly Ala His Ser Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu
              1                      5                      10

Val Arg Pro Ser Gln Thr Leu Ser Leu Thr Cys Thr Ala Ser Gly

```

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	15		20		25	
Ph	Asn Ile Lys Asp Thr Tyr Met	His Trp Val Arg Gln Pr	Pro			
	30	35	40			
Gly Arg Gly Leu Glu Trp Ile Gly Arg	Ile Asp Pro Ala Ser Gly					
	45	50	55			
Asp Thr Lys Tyr Asp Pro Lys Phe Gln	Val Arg Val Thr Met Leu					
	60	65	70			
Val Asp Thr Ser Ser Asn Gln Phe Ser	Leu Arg Leu Ser Ser Val					
	75	80	85			
Thr Ala Ala Asp Thr Ala Val Tyr Tyr	Cys Ala Asp Gly Met Trp					
	90	95	100			
Val Ser Thr Gly Tyr Ala Leu Asp Phe	Trp Gly Gln Gly Thr Thr					
	105	110	115			
Val Thr Val Ser Ser Gly Glu Ser						
	120					

(2) INFORMATION FOR SEQ ID NO:7:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 386 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (ix) FEATURE:
 - (A) NAME/KEY: sig_peptide
 - (B) LOCATION: 1-57
- (ix) FEATURE:
 - (A) NAME/KEY: mat_peptide
 - (B) LOCATION: 58-386
- (ix) EATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 1-386
- (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 1
 - (D) OTHER INFORMATION: /note= "pBAG198 insert: VK2 (SVMDY) light chain variable region"

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

ATG	GGT	TGG	TCC	TGC	ATC	ATC	CTG	TTC	CTG	GTT	GCT	ACC	GCT	ACC	45
M t	Gly	Trp	Ser	Cys	Ile	Ile	Leu	Phe	Leu	Val	Ala	Thr	Ala	Thr	
			-15						-10					-5	
GGT	GTC	CAC	TCC	AGC	ATC	GTG	ATG	ACC	CAG	AGC	CCA	AGC	AGC	CTG	90
Gly	Val	His	Ser	Ser	Ile	Val	Met	Thr	Gln	Ser	Pro	Ser	Ser	Leu	
			1					5					10		
AGC	GCC	AGC	GTG	GGT	GAC	AGA	GTG	ACC	ATC	ACC	TGT	AAG	GCC	AGT	135
Ser	Ala	Ser	Val	Gly	Asp	Arg	Val	Thr	Ile	Thr	Cys	Lys	Ala	Ser	
			15					20					25		
CAG	AGT	GTG	ACT	AAT	GAT	GTA	GCT	TGG	TAC	CAG	CAG	AAG	CCA	GGT	180
Gln	Ser	Val	Thr	Asn	Asp	Val	Ala	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	
			30					35					40		
AAG	GCT	CCA	AAG	CTG	CTG	ATC	TAC	TAT	GCA	TCC	AAT	CGC	TAC	ACT	225
Lys	Ala	Pro	Lys	Leu	Leu	Ile	Tyr	Tyr	Ala	Ser	Asn	Arg	Tyr	Thr	
			45					50					55		
GGT	GTG	CCA	GAT	AGA	TTC	AGC	GGT	AGC	GGT	TAT	GGT	ACC	GAC	TTC	270
Gly	Val	Pro	Asp	Arg	Phe	Ser	Gly	Ser	Gly	Tyr	Gly	Thr	Asp	Phe	
			60					65					70		
ACC	TTC	ACC	ATC	AGC	AGC	CTC	CAG	CCA	GAG	GAC	ATC	GCC	ACC	TAC	315
Thr	Phe	Thr	Ile	Ser	Ser	Leu	Gln	Pro	Glu	Asp	Ile	Ala	Thr	Tyr	
			75					80					85		
TAC	TGC	CAG	CAG	GAT	TAT	AGC	TCT	CCG	TAC	ACG	TTC	GGC	CAA	GGG	360
Tyr	Cys	Gln	Gln	Asp	Tyr	Ser	Ser	Pro	Tyr	Thr	Phe	Gly	Gln	Gly	
			90					95					100		
ACC	AAG	GTG	GAA	ATC	AAA	CGT	AAG	TG							386
Thr	Lys	Val	Glu	Ile	Lys	Arg	Lys								
			105												

(2) INFORMATION FOR SEQ ID NO:8:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 128 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Met	Gly	Trp	Ser	Cys	Ile	Ile	Leu	Phe	Leu	Val	Ala	Thr	Ala	Thr	
				-15					-10					-5	

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Gly Val His Ser Ser Ile Val Met Thr Gln S r Pro Ser Ser Leu
 1 5 10
 S r Ala Ser Val Gly Asp Arg Val Thr Il Thr Cys Lys Ala Ser
 15 20 25
 Gln Ser Val Thr Asn Asp Val Ala Trp Tyr Gln Gln Lys Pro Gly
 30 35 40
 Lys Ala Pro Lys Leu Leu Ile Tyr Tyr Ala Ser Asn Arg Tyr Thr
 45 50 55
 Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Tyr Gly Thr Asp Phe
 60 65 70
 Thr Phe Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr
 75 80 85
 Tyr Cys Gln Gln Asp Tyr Ser Ser Pro Tyr Thr Phe Gly Gln Gly
 90 95 100
 Thr Lys Val Glu Ile Lys Arg Lys
 105

(2) INFORMATION FOR SEQ ID NO:9:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 1348 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(ix) FEATURE:

- (A) NAME/KEY: VCAM-1 gene segment
- (B) LOCATION: 1-219
- (D) OTHER INFORMATION: This portion of the sequence corresponds, in part, to Exons I, II and III nucleotide sequence of the VCAM-1 gene of Cybulsky et al. Proc. Nat'l. Acad. Sci. USA 88:7861 (1991).

(ix) FEATURE:

- (A) NAME/KEY: Hinge region
- (B) LOCATION: 220-229
- (D) OTHER INFORMATION: This portion of the sequence corresponds, in part, to Fig. 12A in PCT/US92/0250 and represents the hinge region of Human IgG1 heavy chain constant region.

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- (ix) FEATURE:
 (A) NAME/KEY: Heavy chain constant region 2
 (B) LOCATION: 230-338
 (D) OTHER INFORMATION: This portion of the sequence corresponds, in part, to Fig. 12A in PCT/US92/02050 and represents the heavy chain constant region 2 of Human IgG1 heavy chain constant region.

- (ix) FEATURE:
 (A) NAME/KEY: heavy chain constant region 3
 (B) LOCATION: 339-446
 (D) OTHER INFORMATION: This portion of the sequence corresponds, in part, to Fig. 12A in PCT/US92/02050 and represents the heavy chain constant region 3 of Human IgG1 heavy chain constant region.

ATG	CCT	GGG	AAG	ATG	GTC	GTG	ATC	CTT	GGA	GCC	TCA	AAT	ATA	CTT	45
Met	Pro	Gly	Lys	Met	Val	Val	Ile	Leu	Gly	Ala	Ser	Asn	Ile	Leu	
				5					10					15	
TGG	ATA	ATG	TTT	GCA	GCT	TCT	CAA	GCT	TTT	AAA	ATC	GAG	ACC	ACC	90
Trp	Ile	Met	Phe	Ala	Ala	Ser	Gln	Ala	Phe	Lys	Ile	Glu	Thr	Thr	
				20					25					30	
CCA	GAA	TCT	AGA	TAT	CTT	GCT	CAG	ATT	GGT	GAC	TCC	GTC	TCA	TTG	135
Pro	Glu	Ser	Arg	Tyr	Leu	Ala	Gln	Ile	Gly	Asp	Ser	Val	Ser	Leu	
				35					40					45	
ACT	TGC	AGC	ACC	ACA	GGC	TGT	GAG	TCC	CCA	TTT	TTC	TCT	TGG	AGA	180
Thr	Cys	Ser	Thr	Thr	Gly	Cys	Glu	Ser	Pro	Phe	Phe	Ser	Trp	Arg	
				50					55					60	
ACC	CAG	ATA	GAT	AGT	CCA	CTG	AAT	GGG	AAG	GTG	ACG	AAT	GAG	GGG	225
Thr	Gln	Ile	Asp	Ser	Pro	Leu	Asn	Gly	Lys	Val	Thr	Asn	Glu	Gly	
				65					70					75	
ACC	ACA	TCT	ACG	CTG	ACA	ATG	AAT	CCT	GTT	AGT	TTT	GGG	AAC	GAA	270
Thr	Thr	Ser	Thr	Leu	Thr	Met	Asn	Pro	Val	Ser	Phe	Gly	Asn	Glu	
				80					85					90	
CAC	TCT	TAC	CTG	TGC	ACA	GCA	ACT	TGT	GAA	TCT	AGG	AAA	TTG	GAA	315
His	Ser	Tyr	Leu	Cys	Thr	Ala	Thr	Cys	Glu	Ser	Arg	Lys	Leu	Glu	
				95					100					105	
AAA	GGA	ATC	CAG	GTG	GAG	ATC	TAC	TCT	TTT	CCT	AAG	GAT	CCA	GAG	360
Lys	Gly	Ile	Gln	Val	Glu	Ile	Tyr	Ser	Phe	Pro	Lys	Asp	Pro	Glu	
				110					115					120	
ATT	CAT	TTG	AGT	GGC	CCT	CTG	GAG	GCT	GGG	AAG	CCG	ATC	ACA	GTC	405
Ile	His	Leu	Ser	Gly	Pro	Leu	Glu	Ala	Gly	Lys	Pro	Ile	Thr	Val	

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125										130					135				
AAG	TGT	TCA	GTT	GCT	GAT	GTA	TAC	CCA	TTT	GAC	AGG	CTG	GAG	ATA	450				
Lys	Cys	Ser	Val	Ala	Asp	Val	Tyr	Pro	Ph	Asp	Arg	Leu	Glu	Ile	150				
				140					145										
GAC	TTA	CTG	AAA	GGA	GAT	CAT	CTC	ATG	AAG	AGT	CAG	GAA	TTT	CTG	495				
Asp	Leu	Leu	Lys	Gly	Asp	His	Leu	Met	Lys	Ser	Gln	Glu	Phe	Leu	165				
				155					160										
GAG	GAT	GCA	GAC	AGG	AAG	TCC	CTG	GAA	ACC	AAG	AGT	TTG	GAA	GTA	540				
Glu	Asp	Ala	Asp	Arg	Lys	Ser	Leu	Glu	Thr	Lys	Ser	Leu	Glu	Val	180				
				170					175										
ACC	TTT	ACT	CCT	GTC	ATT	GAG	GAT	ATT	GGA	AAA	GTT	CTT	GTT	TGC	585				
Thr	Phe	Thr	Pro	Val	Ile	Glu	Asp	Ile	Gly	Lys	Val	Leu	Val	Cys	195				
				185					190										
CGA	GCT	AAA	TTA	CAC	ATT	GAT	GAA	ATG	GAT	TCT	GTG	CCC	ACA	GTA	630				
Arg	Ala	Lys	Leu	His	Ile	Asp	Glu	Met	Asp	Ser	Val	Pro	Thr	Val	210				
				200					205										
AGG	CAG	GCT	GTA	AAA	GAA	TTG	CAA	GTC	GAC	AAA	ACT	CAC	ACA	TGC	675				
Arg	Gln	Ala	Val	Lys	Glu	Leu	Gln	Val	Asp	Lys	Thr	His	Thr	Cys	225				
				215					220										
CCA	CCG	TGC	CCA	GCA	CCT	GAA	CTC	CTG	GGG	GGA	CCG	TCA	GTC	TTC	720				
Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser	Val	Phe	240				
				230					235										
CTC	TTC	CCC	CCA	AAA	CCC	AAG	GAC	ACC	CTC	ATG	ATC	TCC	CGG	ACC	765				
Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg	Thr	255				
				245					250										
CCT	GAG	GTC	ACA	TGC	GTG	GTG	GTG	GAC	GTG	AGC	CAC	GAA	GAC	CCT	810				
Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro	270				
				260					265										
GAG	GTC	AAG	TTC	AAC	TGG	TAC	GTG	GAC	GGC	GTG	GAG	GTG	CAT	AAT	855				
Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	285				
				275					280										
GCC	AAG	ACA	AAG	CCG	CGG	GAG	GAG	CAG	TAC	AAC	AGC	ACG	TAC	CGG	900				
Ala	Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	300				
				290					295										
GTG	GTC	AGC	GTC	CTC	ACC	GTC	CTG	CAC	CAG	GAC	TGG	CTG	AAT	GGC	945				
Val	Val	Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	315				
				305					310										
AAG	GAG	TAC	AAG	TGC	AAG	GTC	TCC	AAC	AAA	GCC	CTC	CCA	GCC	CCC	990				
Lys	Glu	Tyr	Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro					

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320										325					330					
ATC	GAG	AAA	ACC	ATC	TCC	AAA	GCC	AAA	GGG	CAG	CCC	CGA	GAA	CCA	1035					
Ile	Glu	Lys	Thr	Ile	Ser	Asp	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro						
				335					340					345						
CAG	GTG	TAC	ACC	CTG	CCC	CCA	TCC	CGG	GAT	GAG	CTG	ACC	AAG	AAC	1080					
Gln	Val	Tyr	Thr	Leu	Pro	Pro	Ser	Arg	Asp	Glu	Leu	Thr	Lys	Asn						
				350					355					360						
CAG	GTC	AGC	CTG	ACC	TGC	CTG	GTC	AAA	GGC	TTC	TAT	CCC	AGC	GAC	1125					
Gln	Val	Ser	Leu	Thr	Cys	Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp						
				365					370					375						
ATC	GCC	GTG	GAG	TGG	GAG	AGC	AAT	GGG	CAG	CCG	GAG	AAC	AAC	TAC	1170					
Ile	Ala	Val	Glu	Trp	Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr						
				380					385					390						
AAG	ACC	ACG	CCT	CCC	GTG	CTG	GAC	TCC	GAC	GGC	TCC	TTC	TTC	CTC	1215					
Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe	Leu						
				395					400					405						
TAC	AGC	AAG	CTC	ACC	GTG	GAC	AAG	AGC	AGG	TGG	CAG	CAG	GGG	AAC	1260					
Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser	Arg	Trp	Gln	Gln	Gly	Asn						
				410					415					420						
GTC	TTC	TCA	TGC	TCC	GTG	ATG	CAT	GAG	GCT	CTG	CAC	AAC	CAC	TAC	1305					
Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala	Leu	His	Asn	His	Tyr						
				425					430					435						
ACG	CAG	AAG	AGC	CTC	TCC	CTG	TCT	CCG	GGT	AAA	TGA	GTG	CGG		1348					
Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys										
				440					445											

(2) INFORMATION FOR SEQ ID NO:10:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 24 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(ix) FEATURE:

- (A) NAME/KEY:
- (B) LOCATION:
- (D) OTHER INFORMATION: This corresponds to Kinase Primer 370-31.

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

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TCGTC GAC AAA ACT CAC ACA TGC C
 Asp Lys Thr His Thr Cys
 1 5

24

(2) INFORMATION FOR SEQ ID NO:11:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 24 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(ix) FEATURE:

- (A) NAME/KEY:
- (B) LOCATION:
- (D) OTHER INFORMATION: This corresponds to Kinase
Primer 370-32.

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

GTAAATGAGT GCGGCGGCCG CCAA

24

(2) INFORMATION FOR SEQ ID NO:12:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 115 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(ix) FEATURE:

- (A) NAME/KEY:
- (B) LOCATION:
- (D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

GCGGCCGCGG TCCAACCACC AATCTCAAAG CTTGGTACCC GGAATTCAG

50

ATCTGCAGCA TGCTCGAGCT CTAGATATCG ATTCCATGGA TCCTCACATC

100

CCAATCCGCG GCCGC

115

(2) INFORMATION FOR SEQ ID NO:13:

- 64 -

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 39 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(ix) FEATURE:
(A) NAME/KEY:
(B) LOCATION:
(D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

GA	GCT	CGA	GGC	GGC	CGC	ACC	ATG	CCT	GGG	AAG	ATG	GTC	GTG	41
							Met	Pro	Gly	Lys	Met	Val	Val	
							1				5			

(2) INFORMATION FOR SEQ ID NO:14:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 23 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(ix) FEATURE:
(A) NAME/KEY:
(B) LOCATION:
(D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

AA	GTC	GAC	TTG	CAA	TTC	TTT	TAC	23
----	-----	-----	-----	-----	-----	-----	-----	----

(2) INFORMATION FOR SEQ ID NO:15:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 14 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(ix) FEATURE:
(A) NAME/KEY:

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(B) LOCATION:
(D) OTHER INFORMATION:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

TCGACGCGGC CGCG

14

CLAIMS:

1. A method for the prevention of insulin dependent (type I) diabetes comprising administering to a prediabetic individual a composition comprising an anti-VLA4 antibody.
2. A method according to claim 1, wherein the anti-VLA4 antibody selected from the group consisting of HP1/2, HP2/1, HP2/4, L25, and P4C2.
3. A method according to claim 1, wherein the anti-VLA4 antibody is HP1/2, or a fragment thereof, capable of binding to VLA4.
4. A method according to claim 1, wherein the anti-VLA4 antibody is a humanized HP1/2 antibody, or a fragment thereof, capable of binding to VLA4.
5. A method according to claim 1, wherein the composition is administered at a dosage so as to provide from about 0.1 to about 10 mg/kg, based on the weight of the prediabetic individual.
6. A method according to claim 1, wherein the composition is administered in an amount effective to coat VLA4-positive cells in the peripheral blood for a period of 1-14 days.
7. A method according to claim 1, wherein the composition is administered in an amount effective to provide a plasma level of antibody in the prediabetic individual of at least 1 μ g/ml.
8. A method according to claim 1, wherein the composition is administered prior to the development of overt diabetes, as measured by a serum glucose level of less than about 250 mg/dL.
9. A method according to claim 1, wherein the prediabetic individual is a human.
10. A method for the treatment of diabetes comprising administering to a mammal with a susceptibility to diabetes, an antibody, a recombinant antibody, a chimeric antibody, fragments of such antibodies, a polypeptide or a small

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molecule capable of binding to the α_4 subunit of VLA4, or combinations of any of the foregoing, in an amount effective to provide inhibition of onset of diabetes.

11. A method according to claim 10, wherein the antibody, polypeptide or molecule is selected from monoclonal antibody HP1/2; Fab, Fab', F(ab')₂ or F(v) fragments of such antibody; soluble VCAM-1 or fibronectin polypeptides; or small molecules that bind to the VCAM-1 or fibronectin binding domain of VLA4.

12. A method according to claim 10, wherein the composition comprises a plurality of anti-VLA4 monoclonal antibodies or VLA4-binding fragments thereof.

13. A method according to claim 10, wherein the composition is administered at a dosage so as to provide from about 0.1 to about 10 mg/kg of antibody, antibody fragment, polypeptide or small molecule, based on the weight of the susceptible mammal.

14. A method according to claim 10, wherein the composition is administered in an amount effective to coat VLA4-positive cells in the peripheral blood for a period of 1-14 days.

15. A method according to claim 10, wherein the composition is administered in an amount effective to provide a plasma level of antibody in the mammal of at least 1 μ g/ml over a period of 1-14 days.

16. A method according to Claim 11, wherein the soluble VCAM-1 polypeptides comprise VCAM 2D-IgG.

17. A pharmaceutical composition effective to provide inhibition of onset of diabetes consisting essentially of a monoclonal antibody recognizing VLA4 in a pharmaceutically acceptable carrier.

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FIG. 1

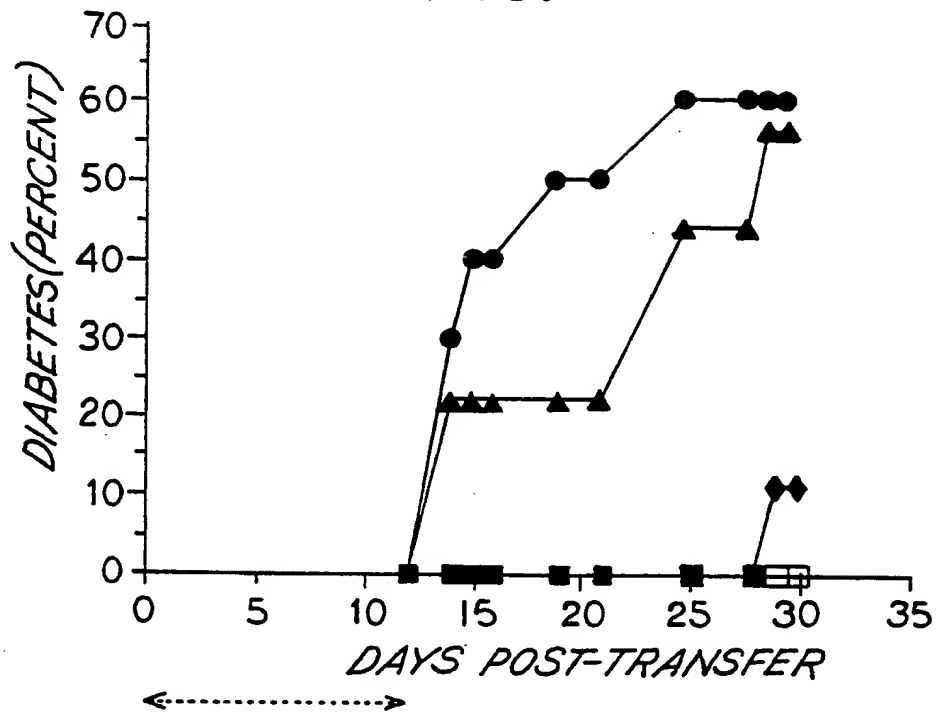
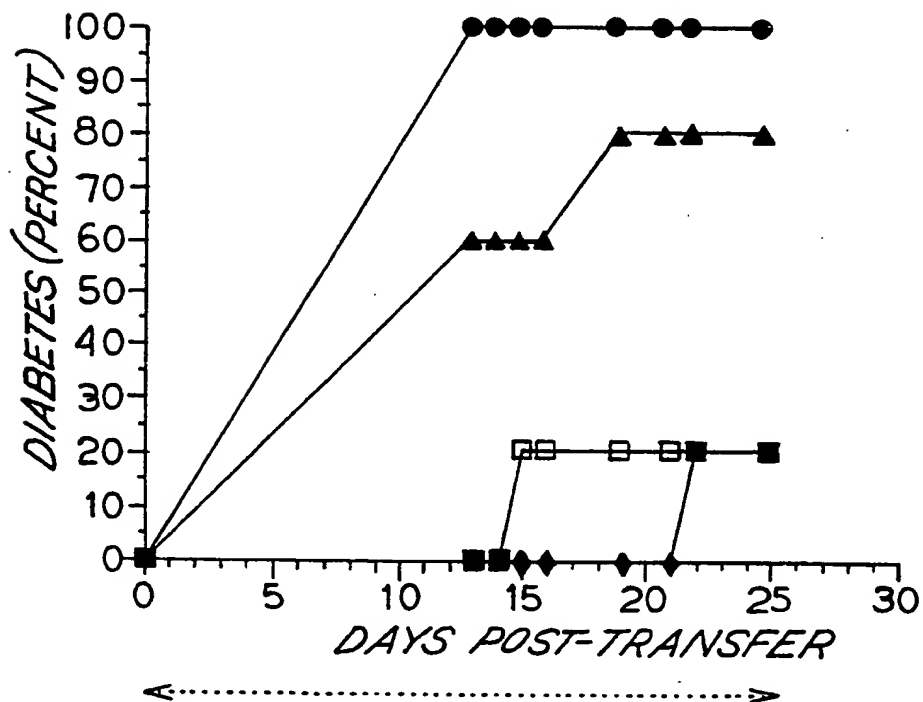


FIG. 2



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FIG. 3

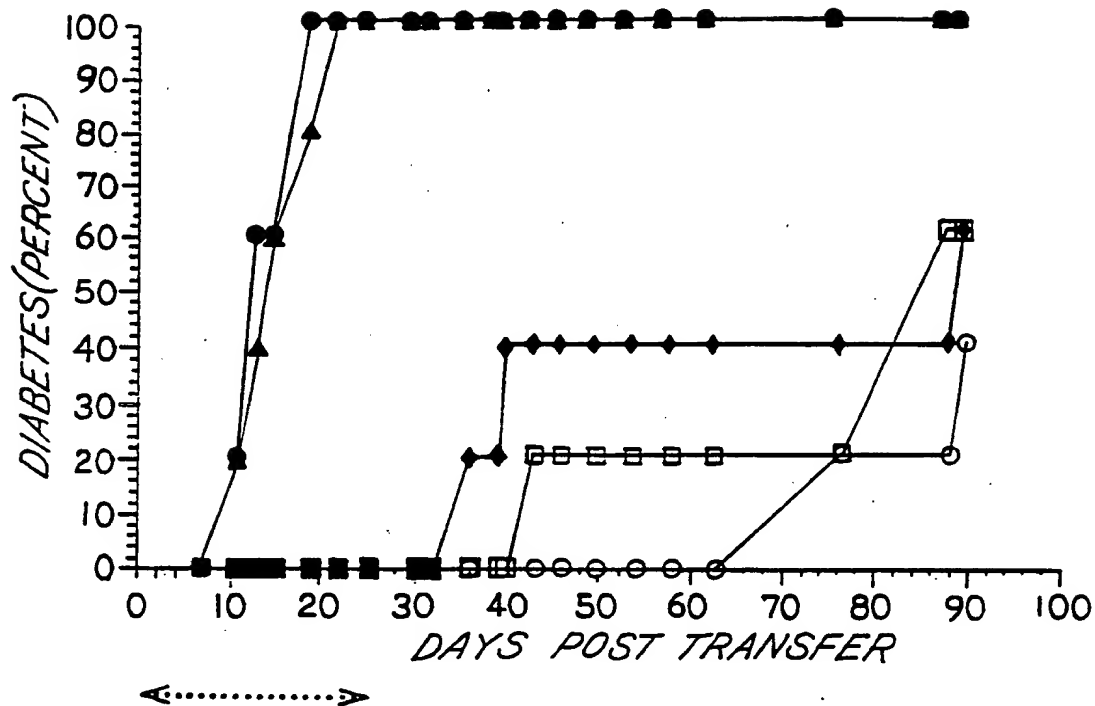
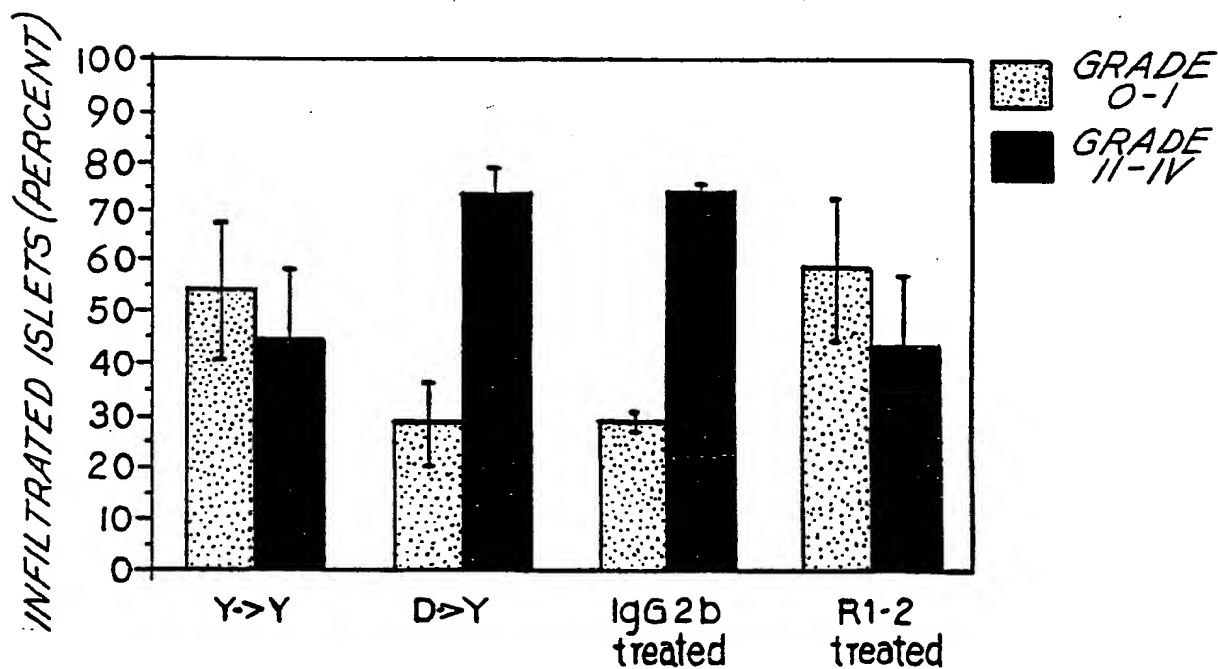
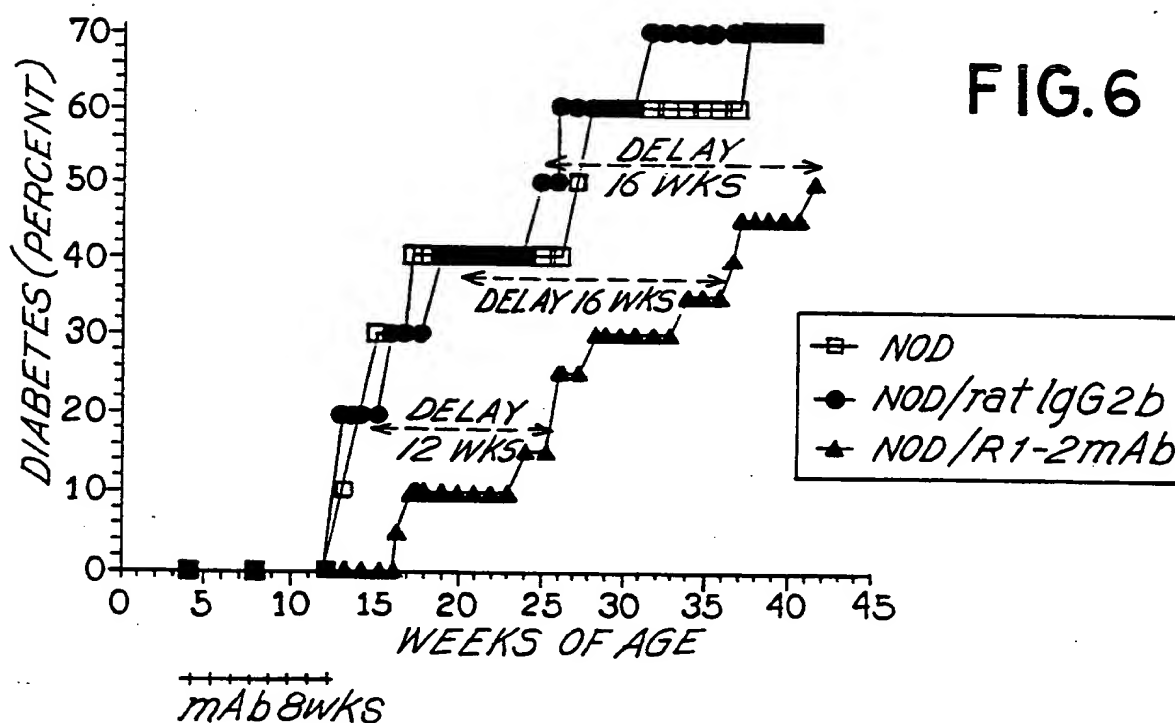
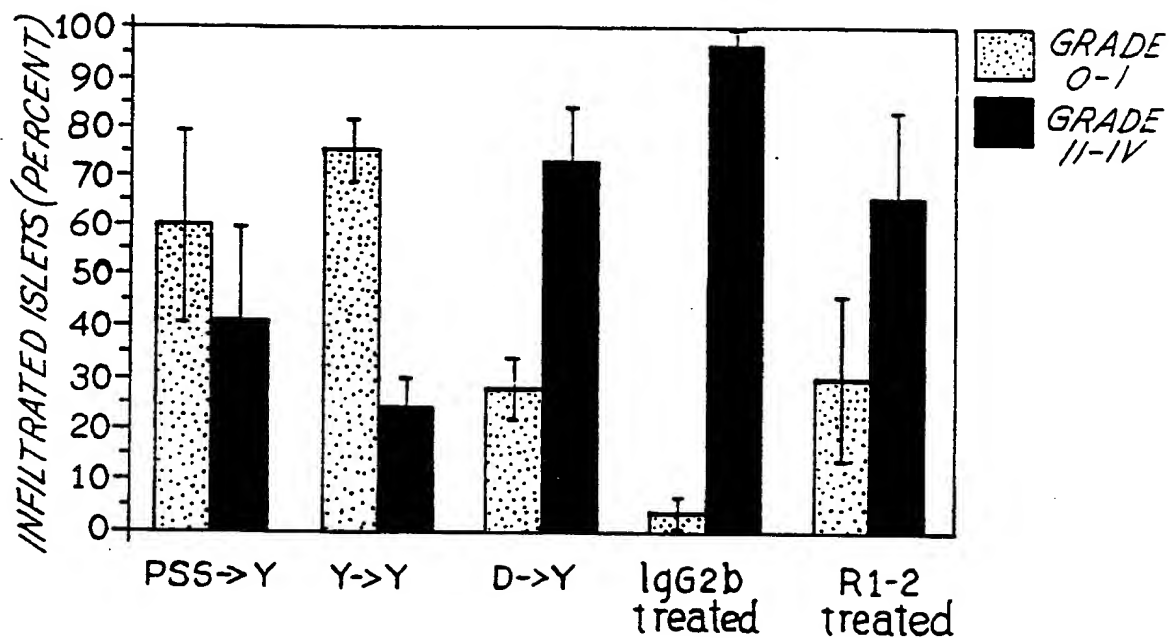


FIG. 4



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FIG. 5



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FIG. 7

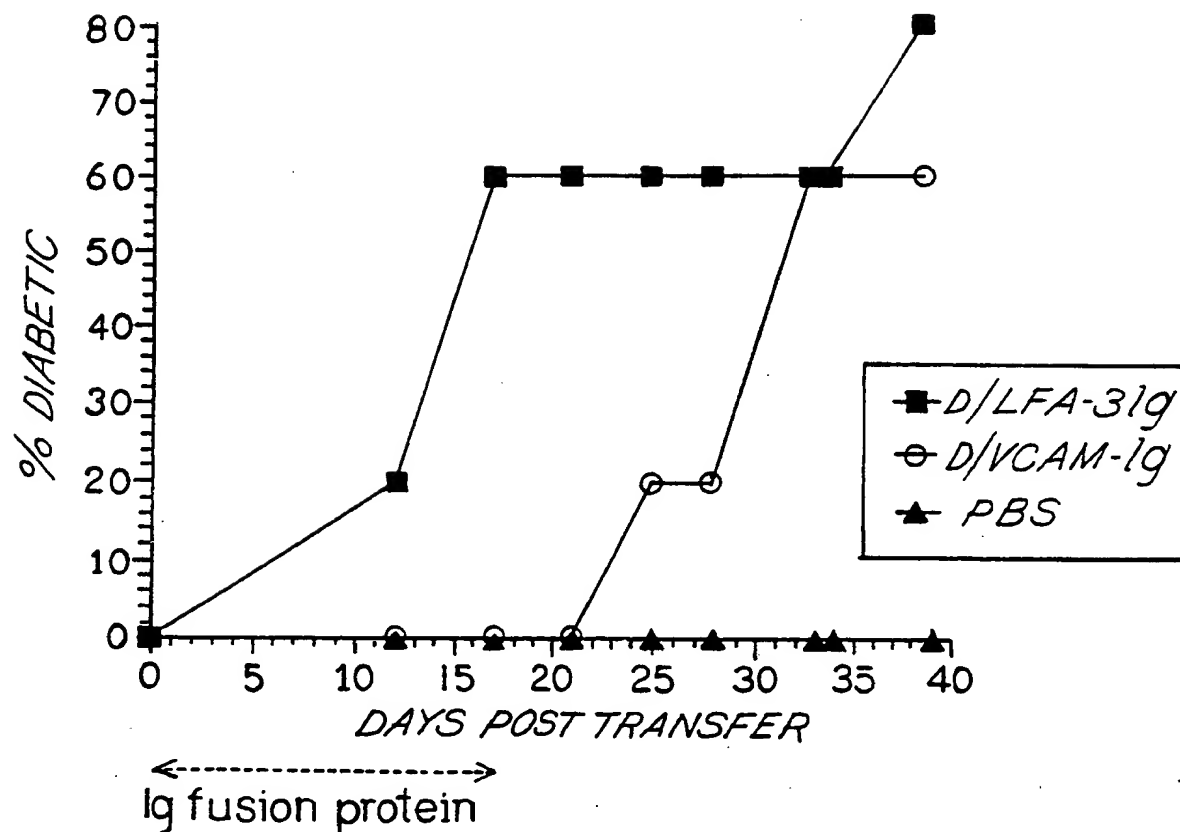


FIG. 8

